

REVIEW

Oligofurostanosides—Furostanol Saponins

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Oligofurostanosides—the furostanol saponins with at least two sugar chains isolated till 1993 from various plants are reviewed along with their physical data. Latest chemical and spectroscopic techniques used for their isolation and characterisation have also been described.

Although a good number of reviews have appeared in various journals covering the spirostanol saponins,¹⁻¹³ but very little efforts have been made to compile the research work done on this new class of glycosides, the oligofurostanosides. The present work is an humble attempt to review all the chemical investigations carried out in this field.

In the recent past the chemistry of natural products has touched new horizons and is rapidly developing as a separate discipline in itself. The importance of this field lies in the fact that on one hand it is the organic chemistry associated with plants, animals, marine and on the other hand it is associated with plant biochemistry and the utility in pharmaceutical chemistry. Phytochemistry, as it is more popularly known, is concerned with enormous types of organic compounds which are not only synthesised by the plants, but are also preserved by them for their growth and protection. Due to this reason phytochemistry involves a study of the structures of chemicals compounds, their biosynthesis and biological functions. The plant kingdom contains enormous organic compounds of different classes, out of which the chemistry of glycosides has been the focal point of research of many modern chemists.

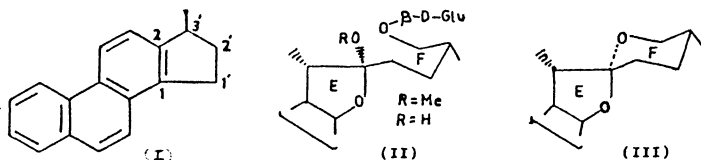
The O-glycosides, which produce copious foam on shaking with water, cause sneezing on inhalation, kill fishes and haemolyse red blood corpuscles are usually termed as 'Saponins' (Latin, *sapo* = soap) and have been classified as: Triterpenoid Saponins, Steroidal-Alkaloidal Saponins and Steroidal Saponins, out of which triterpenoid saponins have been reviewed most abundantly by many workers.

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Steroidal Saponins

These saponins on acidic/enzymatic hydrolysis give C_{27} sapogenins, which possess the carbon ring system of cholesterol along with the remaining 8-carbon atoms as a bicyclic ring system (rings 'E' and 'F'). These saponins on selenium dehydrogenation¹⁴⁻¹⁶ are degraded into Diels' hydrocarbon (3'-methyl 1,2-cyclopentenophenanthrene, (I)).

The C_{27} steroidal aglycones of steroidal saponins have been found to be oxygenated at carbon atoms 16, 22, 26 and carry a hydroxyl group at C_3 . Depending upon the nature of ring 'F', they are termed as furostanolic saponins (open ring 'F' (II)) and Spirostanolic saponins (closed ring 'F', (III)). Sharma *et al.* further¹⁷⁻¹⁹ subclassified them as:



1. Oligofurostanosides and Furostanosides

Furostanolic saponins with ≥ 3 sugars are termed as oligofurostanosides because of their close relationship and resemblance with oligosaccharides on one hand and saponins on the other, whereas the furostanolic saponins with < 3 sugars, not resembling with oligosaccharides, are called as furostanosides.

2. Oligospirostanosides and Spirostanosides

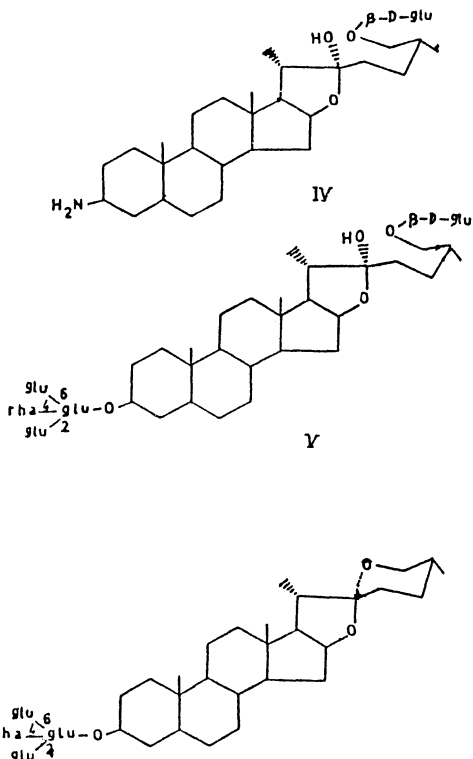
Parallely, the spirostanolic saponins are classified as oligospirostanosides (≥ 3 sugars) and spirostanosides (< 3 sugars).

The first isolation and characterisation of this type of a compound, Jurubine (IV), a furostanolic-alkaloidal saponin, was reported by Schreiber and Ripperger^{20, 21} from *Solanum paniculatum* L. and the very next year came the isolation and characterisation of first such saponin glycoside Sarsaparilloside (V) from *Smilax aristolochiae* Mill. roots by Tschesche *et al.*²², which was termed as Bisdesmoside, with its corresponding oligospirostanoside, Parillin (VI).

These findings were in contrast to Wall's claim²³ but supported Marker's classical suggestion²⁴ that a steroidal saponin is secondarily produced from a steroidal glycoside having an aliphatic chain at C_{17} or via an oligofurostanoside. Oligofurostanosides are, therefore, the real saponin glycosides present in nature and all others are the artefacts, which has been proved by the ageing of the plant material as well as thermal effects^{17-20, 25-27}.

Detection of Oligofurostanosides

Oligofurostanosides can be easily detected by:



Ehrlich reagent test²⁹: The oligofurostanosides on spraying with a solution of *p*-dimethyl aminobenzaldehyde in HCl (Ehrlich reagent) develop a brilliant red colour. The test is specific for pseudocompounds ($\Delta^{20(22)}$ derivative) also. However, furostanolic saponins with 17-OH group are not detected by this reagent.³⁰

Pt. (IV) oxide or NaBH₄ reduction followed by acid hydrolysis^{22, 31}: H₂/Pt oxide or NaBH₄ reduction of oligofurostanosides followed by acid hydrolysis leads mainly to the formation of a dihydroaglycone along with other minor products. Spirostanolic saponins are not affected by this reduction.

IR-spectra^{28, 32-34}: Oligospirostanosides give well defined bands around 860, 900, 920 and 980 cm⁻¹ in IR spectra which are fewer and broader or absent in oligofurostanosides.

Two-dimensional TLC³²: A spot of oligofurostanosides on two-dimensional TLC turns into three diagonal spots because of their continuous conversion into a mixture of two products *viz.* C₂₂-OH and C₂₂-OMe/-OEt in MeOH/EtOH and vice-versa in aqueous medium.

NMR-spectra^{33, 35}: The ¹H-NMR spectra of oligofurostanosides exhibit a characteristic singlet at about δ 3.25 ppm for 22-OMe group.

Crystallisation in MeOH or EtOH^{22, 32}: Oligofurostanosides on crystallisation in MeOH/EtOH yield a mixture of two products *viz.* C₂₂-OH and C₂₂-OMe/-OEt due to continuous transformation of the C₂₂-OH into C₂₂-OMe/-OEt derivatives and vice-versa. However, the oligofurostanosides on

refluxing with water and acetone convert themselves exclusively into C₂₂-OH compounds and on refluxing with dry MeOH into C₂₂-OMe derivatives.

Extraction, Purification and Crystallisation

For the extraction of oligofurostanosides and oligospirostanosides, the air-dried and powdered plant material is first extracted with pet. ether for 6 h usually 3 times to remove oily matter, chlorophyll and any neutral compounds and then with ethyl acetate thrice, 6 h each time, to remove colouring matters etc. Finally, the plant material is subjected to exhaustive methanol/ethanol extraction for 8 h usually four times. The combined alcoholic extract is concentrated under vacuum and examined for their contents with the help of TLC in suitable solvent systems.

The concentrated methanolic/ethanolic extract is dissolved in a minimum quantity of methanol and poured dropwise into large volumes of acetone with constant shaking to precipitate the oligofurostanosides and/or their artefacts. The purification is further achieved through column chromatography using different solvent systems and different adsorbents (*cf.* Table-1).

TABLE-1

Adsorbent	Solvent system	Nature of steroidal saponins
<i>A. Thin layer and column chromatography</i>		
Silica gel	CHCl ₃ : MeOH : H ₂ O 65 : 20-30 : 10	Non-polar ³⁶⁻³⁷
Silica gel	CHCl ₃ : MeOH : H ₂ O 65 : 35 : 10	Neutral and non-polar ³⁸
Silica gel	EtAc : MeOH : H ₂ O 70 : 15 : 15	Acidic and neutral ³⁹
Silica gel	<i>n</i> -BuOH : EtOH : 25% Ammonia sol. 60 : 13 : 30.5	Polar, acidic ⁴⁰
Acidic silica	CHCl ₃ : MeOH : H ₂ O 65 : 35 : 10	Acidic ⁴¹
Alumina (H ₂ O saturated)	Toluene : <i>n</i> -BuOH 2 : 1-1.4	Natural ⁴²
<i>B. Paper and column chromatography</i>		
Paper (formamide impregnated, Whatman No. 1)	CHCl ₃ : THF : C ₅ H ₅ N 10 : 10 : 2 (Formamide saturated)	Neutral ⁴³⁻⁴⁴
Paper (Whatman No. 1)	Iso-BuOH : EtOH : Et ₂ NH : H ₂ O 5 : 5 : 1 : 4	Acidic ⁴⁰
<i>C. Gel chromatography</i>		
Sephadex G-25 or G-50	Water	Saponins of different molecular sizes ⁴⁵⁻⁴⁶

Quantitative separation is generally achieved by column-chromatography,

through elution in the order of increasing number of sugar molecules present in them. Thin layer (TLC) and paper chromatographic (PC) techniques are generally employed to check the purity of these glycosides.

However, small quantities of saponins can also be separated through preparative TLC and PC. Generally, mixed solvents are used for the chromatographic separation of various saponin mixtures. Water saturated solvents afford better separation. The solvent systems containing chloroform, methanol and water in various proportions and silica gel as the adsorbent are most commonly employed.

Two new techniques named as DCCC and HPLC are also frequently used now-a-days for these purposes.

(i) Droplet Counter Current Chromatography

It is a support free separation technique developed by Tumimura *et al.*⁴⁷, employing the passage of droplets of mobile phase through stationary liquid phase and the solute is continuously partitioned between two phases.

The columns are filled with stationary phase and sample after dissolution in selected solvent is charged in to sample chamber. The mobile phase is pumped through sample chamber as a result of which a stream of droplets is formed in immiscible stationary phase. These droplets are then allowed to move through the column, where partitioning of solute occurs in stationary and solute phase, resulting to the separation of components. However, binary systems are not much useful; hence, tertiary and quaternary systems are usually taken up for the preparation of two phases.

This technique has great advantages of taking less time, no loss of sample, separation of small quantity of samples, less requirements for samples, less requirements for sample purification and easy separation of closely resembling compounds and, hence, has achieved significant importance in the isolation, purification and separation of saponins.⁴⁸⁻⁵⁰

(ii) High Performance or High Pressure Liquid Chromatography⁵¹⁻⁵³

A liquid which is a mobile phase in HPLC is propagated through the column by pumps and the sample is introduced at the head of a column through an injector without disturbing the flow of mobile phase as well as column packing. A device called as detector senses measures the sample component from column effluent. The response of the detector is recorded by a recording device. Now-a-days, a number of recording devices like chromatographic data system, integrator, computers can be linked with the HPLC and the results obtained are essentially without errors.

The structure determination of oligofurostanosides involves the following methods:

Enzymatic hydrolysis^{22, 32}: Enzymatic hydrolysis with β -glucosidase cleaves off the β -linked glucose at C₂₆ (in almost all cases with only a few exceptions) leading to the cyclisation of the ring 'F' with C₂₂-OH and formation of oligospirostanoside. If β -D-glucose is the terminal sugar of the sugar chain

attached at C₃ of the aglycone, it will also be liberated leading to the formation of lower oligofurostanosides/furostanosides/oligospirostanosides etc.

Acid hydrolysis⁵⁴⁻⁵⁶: Oligofurostanosides on complete hydrolysis with 8–10% mineral acid yield a steroidal sapogenin and the monosaccharides (sugars) and/or their oxidation products. The genin is identified by mp, mmp, Co-TLC, IR, NMR, MS and also by converting it into acetate and its mp etc. Neutralised aqueous sugar solution after concentration under vacuum at 40–50°C is subjected to descending paper chromatography⁵⁷⁻⁵⁹ on Whatman filter paper No. 1 using the solvent systems, *n*-BuOH : AcOH : H₂O (4 : 1 : 5); EtAc : Py : H₂O (10 : 4 : 3); *n*-BuOH : Py : H₂O (75 : 15 : 10) etc. to identify the sugars. The spots are visualised by spraying with reagents like aniline hydrogen phthalate, *p*-aniline hydrochloride, ammonical silver nitrate solution, etc.

Kiliani hydrolysis⁶⁰: In order to find out the sequence of sugars in the glycone moiety the oligofurostanoside is subjected to hydrolysis with Kiliani mixture (AcOH : H₂O : 35% HCl, 35 : 55 : 10) at room temperature. The sugars liberated with the run of time are studied by PC. The sugars are liberated from the sugar chain(s) starting from the outermost sugar(s) of the chain(s) towards the aglycone. Further, if two or more than two spots are obtained with the run of time, at one particular moment, branching in the sugar moiety is indicated.

Reduction followed by hydrolysis^{22, 31}: Discussed under detection of oligofurostanosides.

CrO₃ oxidation^{22, 33-34, 61}: The form of ring 'F' as well as the nature and number of sugars attached to C₂₆ of the oligofurostanoside can elegantly be proved by CrO₃ oxidation. The oligofurostanoside is converted to peracetate Δ²⁰⁽²²⁾ through acetylation followed by the removal of a molecule of H₂O or MeOH by heating in AcOH, which on oxidation with CrO₃ in AcOH gives an oxidised product. The basic hydrolysis (*t*-BuOH/KOH) of this product affords δ-hydroxy-γ-methyl *n*-valeric acid glucoside, which contains the sugar unit glycosidated at C₂₆ of the sapogenin. A survey of literature showed that so far β-linked D-glucose has only been found to be attached at C₂₆ (a few exceptions). Another product obtained by alkaline hydrolysis is a glycoside containing the main sugar chain(s) of the parent glycoside intact, which on acid hydrolysis yields the corresponding pregnolone and sugars attached at C₃.

Baeyer-Villiger oxidation^{27, 33}: Similar to CrO₃ oxidation, the structure of the side chain in oligofurostanoside can also be confirmed by the Baeyer-Villiger oxidation using H₂O₂-formic acid/acetic acid whereby, 5α, pregnane-3β, 5α, 6β, 16β, 20α pentanol tetraacetate is obtained instead of pregnolone.

Permethylation: The positions of attachment of various sugars with each other as well as with the aglycone are determined by permethylation of oligofurostanoside and methanolysis/hydrolysis studies of the permethylated derivatives. Permethylation is generally carried out by Purdie⁶² (Ag₂O and MeI), Kuhn⁶³ (BaO, Ba(OH)₂ and MeI/DMSO) and Hakomori⁶⁴ (NaH, DMSO and MeI/N₂ atm.) methods, out of which the latter yields better results.

The permethylated saponins on methanolysis in dry MeOH : HCl (5–10% v/v) give a mixture of methyl methylated sugars, which can be identified and estimated by GLC^{34, 61} by comparison of their retention times with authentic samples.

Position(s) of the free hydroxyl group(s) in the methyl methylated sugars show the attachment of different sugars with each other. The methyl methylated sugars can also be identified by their MS⁶⁵⁻⁶⁷ and NMR⁶⁸ studies. The vicinal branching in the sugar moiety can also be determined alternatively through periodate oxidation studies of the methyl methylated sugars, whereby the corresponding methyl methylated sugars disappear. The PC and isolation studies³² of the methylated sugars, obtained by the hydrolysis of the methyl methylated sugars confirm the linkages of different sugars with each other.

Most commonly employed solvent systems for better PC resolution of methylated sugars are: n-BuOH : EtOH : H₂O (5 : 1 : 4)⁵⁹, 2-butanone saturated with 2% NH₃⁶⁹ and Benzene : EtOH : H₂O : ammonia (200 : 47 : 14 : 1).⁷⁰ Aniline hydrogen phthalate is generally used as the developing reagent. Methylated sugars have also been studied by TLC^{44, 71-72} on silica gel and cellulose.

Partial hydrolysis⁷³⁻⁷⁵: An oligofurostanoside containing a large number of sugars is difficult to handle for its glycone moiety's structure determination, since it may involve a good number of branchings. In such cases, the oligofurostanoside is partially hydrolysed by refluxing it with 5% aq. H₂SO₄-MeOH on steam bath for 30 min or 1N-HCl-MeOH on steam bath for 40 min. Usual work up yields the liberated sugars and a mixture of lower glycosides. These new glycosides are then permethylated and subjected to methanolysis followed by hydrolysis. The resulting methyl-methylated sugars/methylated sugars are identified by PC and compared with those obtained from the permethylated oligofurostanoside to fix up the exact linkages between the sugars themselves and with the aglycone.

Molecular rotation: The α - and β -configurations of different sugars in the oligofurostanosides have been proved by molecular rotation values.⁷⁶⁻⁷⁸ For calculating the theoretical values, the aglycone contribution is calculated from the known values of the corresponding methyl sugars. The best befitting theoretical and experimental molecular rotation values give the configuration of the particular sugar.

Methyl sugars	[M] _D
β -methyl-D-glucopyranoside	-66°
α -methyl-D-glucopyranoside	+309°
β -methyl-D-galactopyranoside	+10°
α -methyl-D-galactopyranoside	+380°
β -methyl-D-xylopyranoside	-108°
α -methyl-D-xylopyranoside	+253°
β -methyl-L-arabinopyranoside	+403°
α -methyl-L-arabinopyranoside	+28°
β -methyl-L-rhamnopyranoside	+170°
α -methyl-L-rhamnopyranoside	-111°

ORD and CD: From the sign and shape of the ORD and CD curves, not only the carbonyl groups in the different positions in the steroidal aglycone can be located but also their chemical environment can be traced easily.⁷⁹⁻⁹⁰ Some of the generalisations provided by these studies are as follows:

1. Alternation of A/B ring junction does not affect the rotatory dispersion curve to any appreciable extent.
2. Sapogenins with 22α -configurations with weak chromophores ($-\text{OH}$, isolated $>\text{C}=\text{C}<$) give $-ve$ plain curves and those having 22β -configuration give $+ve$ plain curves.
3. Introduction of carbonyl group at C_3 , C_{11} , C_{12} and C_{13} affects the $-ve$ rotation by a strongly $+ve$ one.
4. Change in the stereochemistry at C_8 and C_{14} does not appreciably affect the shapes of the curves.

UV spectroscopy: This technique has very successfully been used for the detection of ethylenic bond (193–205 nm, depending upon the position and environment of the double bond) and the carbonyl chromophores (λ_{max} and ϵ_{max} values and the well defined additional band at approximately 280 nm) in steroidal compounds.^{91–95}

IR spectroscopy: IR spectra have provided valuable information about the structural features, various functional groups and stereochemistry at various centres of oligofuro- and oligospirostanosides as well as their aglycones^{28, 32–34, 96–105} e.g.

1. The presence of spiroketal system is visualised by the presence of four characteristic bands around 980, 920, 900 and 860 cm^{-1} . In case of (25S) 920 cm^{-1} band is stronger than 900 cm^{-1} and in case of (25R) 900 cm^{-1} band is stronger than 920 cm^{-1} .
2. Open 'F' ring compounds have either no or broader and fewer above characteristic bands but also their bands below 2830 cm^{-1} are shifted to lower frequency.
3. The absence of any one of the spiroketal band usually means the absence of ring 'F'.
4. A band at about 1240 cm^{-1} indicates trans A/B ring junction or Δ^5 unsaturation.
5. The intensities of the typical spiroketal absorptions are reduced to a great extent by the presence of substitution in ring 'F'.

NMR spectroscopy¹⁰⁶: (i) $^1\text{H-NMR}$: The $^1\text{H-NMR}$ study of steroidal sapogenins in deuterio-chloroform is particularly of great help in stereochemical assignments¹⁰⁷, e.g.,

1. In (25R)-methyl series, the equatorial (25R)-methyl protons resonate at a higher field than the C_{20} -methyl protons.
2. The equatorial (25R)-methyl protons resonate at a higher field than the corresponding axial (25R)-methyl protons.

In pyridine solution the spectral differences between (25R)- and (25S)-methyl series become more distinct,¹⁰⁸ further supported by Tori and Aono¹⁰⁹, who also showed the validity of Zurcher's additivity rules¹¹⁰ by studying the effect of substituents on the positions of the proton signals for the four methyl groups.

$^1\text{H-NMR}$ study of oligofurostanosides etc. also throws much light on the structure of the glycoside molecule. The linkages (α - or β -)^{33–35, 111–113} of various sugars in saponins are very precisely given by the signals and coupling constants of the anomeric protons of sugars. The measurements are usually made indirectly,

via the spectra of oligoglycoside permethylates which are soluble in CHCl_3 . A few generalisations are summarised as follows:

1. The anomeric protons of various sugars give peaks in down field region (δ 4.0–6.30).
2. D-sugars generally occur with β -linkages and are characterised by their high coupling constants (J, 6–9 Hz).
3. The rarely occurring α -linkages of corresponding sugars have low coupling constants (J, 2–4 Hz).
4. The coupling constants of the commonly α -linked occurring sugars, L-rhamnose and L-arabinose are J, ca. 2 Hz and J, ca. 6–8 Hz respectively.
5. The furostanolic nature of oligofurostanosides is shown^{33, 35} by 22-OME signals at about δ 3.25 ppm.

(ii) $^{13}\text{C-NMR}$: This technique provides a non-destructive way for the characterisation and identification of an oligofuro- and oligospirostanoside. For structure elucidation of new compounds by $^{13}\text{C-NMR}$ studies, it is always desirable to compare the observed data with the recorded data for model and related compounds. Mahato *et al.*¹¹⁴ utilised $^{13}\text{C-NMR}$ for the structure elucidation of dioscin and gracillin by a comparison of their peaks with those of their aglycone diosgenin and the sugar moieties using known chemical shifts.¹¹⁵ Characteristic chemical shifts are observed for α - and β -positions of the —OH group where glycosidation takes place.

Thakur and Agrawal *et al.*¹¹⁶ have compiled the $^{13}\text{C-NMR}$ chemical shifts of naturally occurring oligofuro- and oligospirostanosides. In general, $^{13}\text{C-NMR}$ spectra are recorded under proton-noise (broad band) decoupling¹¹⁷ in order to avoid signal overlapping. Usually, $^{13}\text{C-NMR}$ shielding is not very sensitive to solvent but as solute-solvent interactions occur; hence change in chemical shifts can be observed. To avoid this change mainly chloroform and pyridine solvents are used.¹¹⁸

$^{13}\text{C-NMR}$ spectral data of the oligofurostanosides differs significantly from that of the oligospirostanosides, particularly in the chemical shifts for carbon atoms of ring 'E' and 'F'. Oligofurostanosides exhibit C_{22} peaks at 110.8 ppm and at 113.5 for $\text{C}_{22}\text{-OH}$ and $\text{C}_{22}\text{-OMe}$ derivative respectively. The $\text{C}_{22}\text{-OMe}$ signals usually occur at 47.2 ± 0.2 ^{119, 120} though in a few cases at 56.5 ppm.¹²¹ Table-2 shows the chemical shifts for the parent steroidal skeleton.¹²² These values show that the chemical shifts for the ring 'A' and 'B' carbon atoms are markedly affected, hence helpful for differentiating 5α , 5β and Δ^5 steroidal sapogenins because the signals for C_5 and C_{19} exhibit more variation.

The number of anomeric signals determine the number of sugars. The matching values of these signals in the same solvent lead to their identification^{123–125} as solvents alter the chemical shift markedly^{126, 127}. Furanose sugars are distinguished by chemical shifts for C_1 , C_2 and C_4 by 4–14 ppm downfield whereas C_5 by 4–7 ppm upfield as compared to the pyranose form^{126, 128}.

A close resemblance of the chemical shifts due to a terminal sugar with respect to a methyl-O-glycoside leads to its characterisation, whereas chemical shifts of inner sugars differ significantly in comparison to methyl-O-glycoside due to α - β effects of the glycosidation¹²⁹. The glycosidation causes a downfield shift of

4.2–8.5 ppm of the α -carbon, the hydroxyl of which has been directly involved in the glycosidation while neighbouring β -carbon atom shows an upfield shift of 0.05–2.0 ppm in oligoglycosides. The upfield shifts of the β -carbon atoms are quite informative but less consistent, whereas the downfield shift of the α -carbon is characteristic enough for the establishment of inter-glycosidic (α , β) linkages.

TABLE-2
 ^{13}C -NMR CHEMICAL SHIFTS

C Atom No.	Furostanosides						Spirostanosides					
	5 α		5 β^*		Δ^5		5 α		5 β		Δ^5	
	22OH	22OMe	22OH	22OMe	22OH	22OMe	25R	25S	25R	25S	25R	25S
1.	38.5	38.5	37.4	37.4	39.9	39.9	38.7	38.7	37.6	37.6	39.9	40.5
2.	22.2	22.2	21.3	21.3	22.4	22.4	22.2	22.2	21.3	21.3	22.4	23.3
3.	27.1	27.1	27.3	27.3	28.0	28.0	26.8	26.8	27.0	27.0	28.0	27.7
4.	29.2	29.2	27.5	27.5	33.0	33.0	29.0	29.0	27.2	27.2	33.0	34.2
5.	47.1	47.1	43.7	43.7	143.7	143.7	47.1	47.1	43.7	43.7	143.7	144.7
6.	29.2	29.2	27.6	27.6	119.0	119.0	29.0	29.0	27.4	27.4	119.0	118.7
7.	32.4	32.4	26.8	26.8	32.0	32.0	32.4	32.4	26.8	26.8	32.0	32.3
8.	35.4	35.4	35.7	35.7	31.4	31.4	35.2	35.2	35.5	35.2	31.4	31.8
9.	54.8	54.8	40.6	40.6	50.1	50.1	54.8	54.8	40.6	40.6	50.1	50.5
10.	36.4	36.4	35.6	35.6	36.6	36.6	36.3	36.3	35.5	35.5	36.6	37.0
11.	21.1	21.1	21.0	21.0	20.9	20.9	20.7	20.7	20.6	20.6	20.9	21.2
12.	40.2	40.2	40.3	40.1	39.1	39.8	40.2	40.0	40.3	39.9	39.1	40.0
13.	41.2	41.2	41.2	41.3	41.2	41.2	40.6	40.5	40.6	40.7	40.2	40.5
14.	56.2	56.5	56.5	56.5	56.2	56.2	56.5	56.2	56.5	56.5	56.5	56.8
15.	32.3	31.4	32.4	31.4	32.3	31.4	31.8	31.7	31.7	31.7	31.8	32.2
16.	81.3	81.0	81.5	81.2	81.3	81.0	80.8	80.8	81.0	80.9	80.7	81.7
17.	63.9	63.8	63.9	63.8	63.9	63.8	62.3	61.9	62.3	62.3	62.1	62.8
18.	16.7	16.5	16.6	16.5	16.7	16.5	16.5	16.5	16.4	16.4	16.3	16.4
19.	23.6	19.6	24.2	19.6	23.6	19.6	12.3	12.4	24.2	23.9	19.7	19.6
20.	40.6	42.0	40.6	42.1	40.6	42.0	41.6	42.1	41.6	42.2	41.6	42.5
21.	16.4	16.5	16.4	16.5	16.4	16.5	14.5	14.3	14.5	14.3	14.5	14.9
22.	110.7	112.5	110.9	112.7	110.7	112.5	109.0	109.5	109.2	109.7	109.9	109.7
23.	30.2	30.3	30.2	30.3	30.2	30.3	31.4	27.0	31.4	27.1	31.4	27.6
24.	29.9	28.0	29.8	28.0	29.9	28.0	28.9	25.9	28.8	25.8	28.8	26.2
25.	29.9	35.2	29.9	35.4	29.9	35.2	30.3	25.8	30.3	26.0	30.3	26.4
26.	68.1	67.6	68.2	67.8	68.1	67.6	66.7	65.0	66.8	65.2	66.7	65.1
27.	14.4	17.4	17.4	17.4	17.4	17.4	17.1	16.0	17.1	16.1	17.1	16.0
OMe	—	48.9	—	48.9	—	48.9	—	—	—	—	—	—

*Values derived by comparisons of the data in various references.

^{13}C -NMR of pyranoside reveals that the coupling constant for the anomeric carbon atom strictly depends upon the orientation of anomeric hydrogen. The one bond coupling constants for C_2 to C_6 of sugars vary in the range of 142–148 Hz while the anomeric carbon exhibits a larger value of 160–175 Hz^{130–131}. For pyranose with an axial H_1 (β -anomer, 160 Hz) the value is about 10 Hz lower than the corresponding value in equatorial H_1 (α -anomer, 170 Hz).

The point of attachment of sugar with the aglycone leads to the downfield shift of the α -carbon atom and upfield shift of adjacent carbon atom.¹³²⁻¹³⁶ In most of the cases (C_3 -sugar chain) it is 6.6 ± 1.0 ppm downfield shift with lower shielding of C_2 (1.1–3.0 ppm) than C_4 (1.8–4.6 ppm). In oligofurostanosides, in most of the cases having D-glucose at C_{26} , a 6.8 ± 0.3 ppm downfield shift along with usual 1.8 ± 0.4 ppm upfield shift of C_{25} has been observed. Table-3 shows the ^{13}C -NMR chemical shifts and J_{CH} values (in parentheses) of methyl glycopyranosides which commonly occur in oligofurostanosides (D_2O as a solvent). The application and utility of this technique can be exemplified by the structure elucidation of chloromaloside-A and chloromaloside-B¹³⁷.

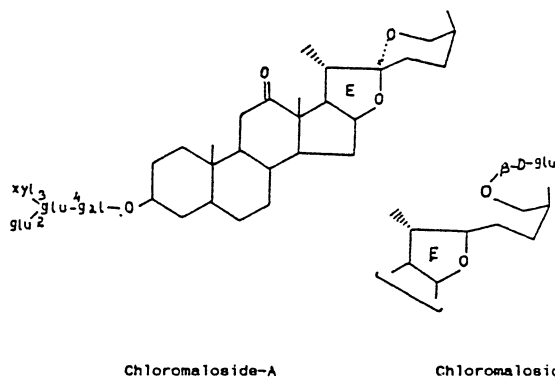


TABLE-3

Methylglycoside	C_1	C_2	C_3	C_4	C_5	C_6
β -D-glucopyranoside	103.7 (160)	73.7 (145)	75.5 (143)	70.3 (141)	75.5 (143)	61.7 (144)
α -D-glucopyranoside	99.9 (170)	72.2 (148)	73.9 (147)	70.4 (145)	71.9 (146)	61.5 (145)
β -D-galactopyranoside	104.1 (160)	71.2 (146)	73.3 (141)	69.1 (146)	75.3 (141)	61.4 (144)
α -D-galactopyranoside	99.8 (170)	69.9 (146)	70.2 (145)	68.9 (146)	71.2 (143)	61.8 (143)
β -D-xylopyranoside	104.8 (159)	73.9 (144)	76.7 (144)	70.1 (147)	66.0 (150)	—
α -D-xylopyranoside	100.3 (170)	72.3 (146)	74.2 (145)	70.3 (146)	61.9 (148)	—
β -L-arabinopyranoside	100.7 (169)	69.8 (145)	69.8 (145)	69.1 (145)	63.4 (149)	—
α -L-arabinopyranoside	104.7 (160)	71.6 (148)	73.3 (143)	69.1 (146)	66.9 (150)	—
β -L-rhamnopyranoside	105.7	72.2	75.4	73.8	73.5	18.5
α -L-rhamnopyranoside	102.4	71.9	72.5	73.6	69.4	18.4

^{13}C -NMR CHEMICAL SHIFTS OF SUGAR MOIETIES IN Py

Sugars	Carbons	Chloromaloside-A	Chloromaloside-B
Galactose	1	102.4	102.5
	2	73.1	73.1
	3	75.5	75.6
	4	79.8	79.9
	5	76.0	76.1
	6	60.7	60.7
Glucose	1	104.8	105.6
	2	81.2	82.1
	3	87.0	87.2
	4	74.4	70.4
	5	77.4	77.5
	6	62.4	62.5
Glucose	1	104.8	105.0
	2	75.0	75.2
	3	78.5	78.6
	4	71.1	71.1
	5	77.7	77.8
	6	62.9	62.9
Xylose	1	104.6	104.7
	2	75.3	75.3
	3	78.5	78.6
	4	70.7	70.7
	5	67.3	67.3
26-O-Glucose	1	—	105.0
	2	—	75.0
	3	—	78.5
	4	—	71.8
	5	—	78.4
	6	—	62.9

 ^{13}C -NMR CHEMICAL SHIFTS OF AGLYCONO MOIETIES IN Py

Carbons	Chloromaloside-A	Chloromaloside-B
1.	36.6	36.7
2.	29.6	29.7
3.	77.3	77.3
4.	34.6	34.7
5.	44.5	44.5
6.	28.6	28.6
7.	31.4	31.4
8.	34.3	34.4
9.	55.6	55.7
10.	36.3	36.3

Carbons	Chloromaloside-A	Chloromaloside-B
11.	38.0	37.9
12.	212.9	212.8
13.	55.3	55.5
14.	55.9	55.9
15.	31.7	31.7
16.	79.8	79.8
17.	54.1	55.3
18.	16.3	16.0
19.	11.7	11.7
20.	43.1	41.1
21.	13.7	15.0
22.	109.8	112.8
23.	26.4	30.9
24.	26.1	28.2
25.	27.5	34.4
26.	65.2	74.6
27.	16.1	17.5
OMe	—	47.4

2D-NMR¹⁰⁶: Two-dimensional NMR, now-a-days, has become a major tool to determine the sequence of different sugar molecules in various glycosides by the following two techniques:

(a) COSY: Homonuclear correlation spectroscopy

(b) NOESY: Heteronuclear correlation spectroscopy

COSY¹³⁸⁻¹⁴¹ reveals the interpretation connectivities and hence identifies the location of the branching points of each sugar in the glycoside molecule. Counting of protons and determining the coupling constants reveal the nature of individual sugars. Usually, 1-D-normal spectrum, 2-D-COSY and 2-D long range COSY are sufficient to identify the sequence of the sugars in reasonably complex steroidal saponins.¹⁴² The observation of typical splitting patterns for characteristic protons H(5) of glucose, rhamnose, arabinose and xylose provides a short cut for the identification of sugars.

NOESY spectrum¹⁴³ reveals the Nuclear Overhauser Effect [NOE] between intra- and inter-protons of sugar molecules, which makes the identification of monosaccharides and their sequencing even more easy.

Mass spectrometry: Mass spectrometry is an important technique for establishing the structure of saponinins and their glycosides. All techniques *i.e.*, EIMS, FDMS, D/CIMS, FABMS and Unimolecular MIKE have been utilised for this purpose with great success.

(a) *EIMS*: EIMS not only furnishes the correct molecular weight but also indicates the positions of certain substituents in saponinins by comparing the characteristic ion peaks with those of unsubstituted standard saponinins¹⁴⁴⁻¹⁵¹.

The characteristic fragmentation pattern of these compounds differentiates them from other classes of compounds. Substituents in the molecule induce a shift in the ion peaks according to their molecular weights. Stereochemical changes in the skeleton cause significant changes in mass fragmentation.

EIMS of permethylated oligofurostanosides and oligospirostanosides^{33, 151} also impart invaluable information about the sequence of sugars in the glycone part through the characteristic ion peaks as:

m/z	Sugars
219, 187	D-Glucose
189, 157	L-Rhamnose
175, 143	L-Arabinose or D-Xylose
331	D-Glucose peracetate
273	L-Rhamnose peracetate
259	D-Xylose peracetate

(b) *FDMS*: To overcome the difficulties in EIMS like, derivatisation of glycoside, evaporation, thermal excitation and unsuitability for substances of low volatility, extensive fragmentation with small or undetectable molecule ion abundances and no molecular ion peaks from oligofuro- and oligospirostanosides, modern techniques like CI (Chemical Ionisation)¹⁵²⁻¹⁵⁴ and FI (Field Ionisation)¹⁵⁵ have been used. However, the thermal stress required for evaporation can't be avoided with these methods which may cause complete decomposition of the sample prior to ionisation.

*FDMS*¹⁵⁶, although only a recent offspring of Field Ionisation mass spectrometry has been found to be highly useful for the structure elucidation of non-volatile and thermally labile substances. In this method, the sample is absorbed on the emitter in a solid layer and is then ionised in the absorbed state. *FDMS* gives important information about the sequence of the sugar moieties present in the glycoside with only a few peaks in comparison to those present in the EIMS spectrum and hence it is easy to interpret.

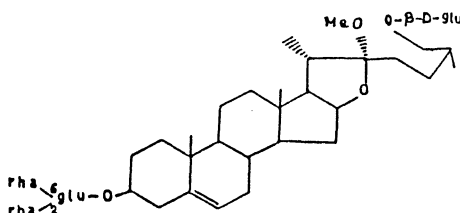
Inorganic impurities, mostly alkali halides, present in the plant glycosides, play an important role in the formation of ions. After a detailed study of *FDMS* of a series of oligofuro- and oligospirostanosides¹⁵⁷⁻¹⁶², the following generalisations have been drawn:

- (i) Alkali metal ions cationisation leads to the formation of most intense $[M + Na]^+$ and less intense $[M + K]^+$ peaks.
- (ii) Sometimes less intense $[M + 2Na]^{2+}$ ions are also recorded.
- (iii) $[M]^+$ and $[M + H]^+$ ions, though formed, are very uncommon.
- (iv) The molecular ion peak for the loss of terminal sugar is also observed.
- (v) Simultaneous losses of two sugars from different positions of branched glycone moiety of oligofuro- and oligospirostanosides are observed as less intense peaks.
- (vi) Cleavage of sugar units at glycosidic oxygen has been explained by analogy with well established mechanism of solvolysis in solution chemistry. The $[H]^+$ ion may be generated by chemically induced reactions.

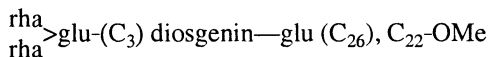


(vii) The cleavage of O-glycoside bond between the aglycone and the sugar attached to it is observed.^{157, 159}

The application and utility of this technique can easily be exemplified by the structure elucidation of Sprengeroside-B¹⁹ as follows:



- m/z 1069 $[(M + K)\text{-MeOH}]^+$, 1053 $[(M + \text{Na})\text{-MeOH}]^+$ (base peak) and 538 $[(M\text{-MeOH}) + 2\text{Na}]^{2+}$ not only confirm the purity but also prove the presence of C_{22} methoxy group.
- The molecular ion peak at m/z 1062 is also observed.
- The peaks at m/z 939 $[(M + \text{Na})\text{-}146]^+$ and 793 $[(M + \text{Na})\text{-}(146 + 146)]^+$ confirm the presence of two terminal rhamnose units.
- The peaks at m/z 923 $[(M + \text{Na})\text{-}162]^+$ and 777 $[(M + \text{Na})\text{-}(162 + 146)]^+$ prove the presence of terminal glucose and a terminal rhamnose unit..
- The peaks displayed at m/z 613 $[(M + \text{Na} + \text{H}_2\text{O})\text{-}454]^+$ confirm the complete sugar chain at C_3 of aglycone.
- The results clearly establish the structure as:



(c) *D/CIMS*: As compared to few informations obtained from FDMS the new technique *D/CIMS*¹⁶³⁻¹⁶⁴ is more informative. The measurements are carried out on a quadrupole MS equipment with ammonia feedline. The emitter is a coiled tungsten wire where the solubilised sample is applied. Ammonia is generally used as the reactant gas since it provides a softer ionisation than methane or isobutane and thus gives invaluable informations. The main advantages of this technique over FDMS and EIMS are:

- Exact molecular weight of glycoside is obtained.
- No derivatisation of glycoside is required.
- Sequence of sugars is also established.
- The characteristic fragmentations of saponins are also obtained which are not predominantly present in FDMS.

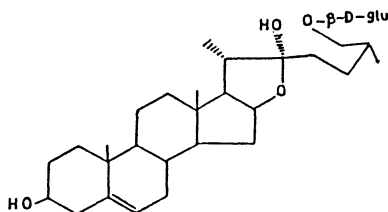
(d) *FABMS*: Barber *et al.*¹⁶⁵ invented the technique of *FABMS* in 1981 to overcome some of the problems of *FDMS* as it was difficult in practice and the ions providing molecular weight information were produced only transiently. In *FABMS* an ion surface is developed to accommodate the solid materials and the phenomenon of ion sputtering employing a beam of fast neutral atom, especially of Ar if 2–8 KeV as the primary particle is used. The sample is normally dissolved in glycerol/*m*-nitrobenzyl alcohol (matrix) to facilitate the production of sample ions in high abundance for relatively long periods. Molecular weight information is usually obtained from $[M + H]^+$ ions in +ve ion spectrum and from $[M - H]^-$ ion in -ve ion spectrum. Odd-electron molecular ions are not normally produced in abundance which give structural informations. Barber *et al.* believed that the $[M + H]^+$ and $[M - H]^-$ ions are either formed by proton transfer reactions which may occur as the molecules are bombarded and pass into gas phase or those already existing in matrix.

FABMS exhibit protonated or sodium adduct¹⁶⁶ molecular ions, which give a series of characteristic fragment ions promising to give a clear picture for defining the sequence of constituents of sugar moieties of the glycosides and also for identifying the glycone (sugars).

Sharma *et al.*¹⁹ have for the first time successfully used this technique for structure elucidation of oligofurostanosides. *FABMS* has been found to have some specific advantages, *viz.*,

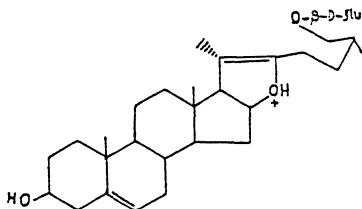
- (i) Mass spectra of molecules with relatively high molecular weight can easily be obtained.
- (ii) Volatilisation of sample is not required and so no thermal effects are observed.
- (iii) No sample derivatisation is required.
- (iv) The method works in either polarity and gives pseudomolecular ion sensitivity with structurally significant fragmentation.
- (v) Ionisation occurs from the solid at room temperature.

The application and utility of this technique can be explained by the structure determination of Sprengerose-E.¹⁹



Sprengerose-E

- (a) Molecular ion peak m/z 595 $[M + H]^+$ was observed.
- (b) The peak at m/z 397 $[(M + H) - (162 + 2H_2O)]^+$ suggested the presence of one terminal glucose and two free —OH groups.
- (c) The peak at m/z 577 $[(M + H) - H_2O]^+$ seems to be observed due to the formation of the following fragment:



All these results clearly established the structure of Sprengerose-E as given above.

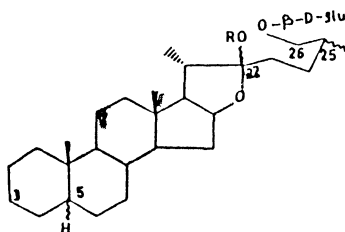
(E) *Unimolecular-mike method*¹⁶⁷: The distinction between A/B and C/D *cis* and *trans* isomers is usually not possible by normal EIMS, which consists of reaction products formed in ion source from high energy reactions in a few microseconds or less. However, the possibility of more stereochemical information might be forthcoming for lower energy reactions, occurring in Unimolecular Mass Analysed Ion Kinetic Energy (MIKE) Spectrometry. This is because

- (i) The spectra are simpler, being only of the reaction products of one ion, *e.g.*, molecular ion.
- (ii) The occurrence of unimolecular reactions of metastable ion is very sensitive to change in the critical energies of these reactions.

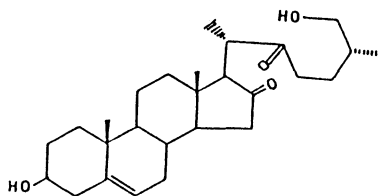
By the study of MIKE spectra of various steroidal compounds, the following generalisations have been carried out:

- (a) For *cis* A/B ring junction (5β) the change in critical energy for reaction involving C_{19} methyl group loss is less as compared to *trans* A/B ring junction (5α).
- (b) For *cis* C/D ring junction (14β) the change in critical energy for the reaction involving C_{18} methyl group loss is greater as compared to *trans* C/D ring junction (14α).

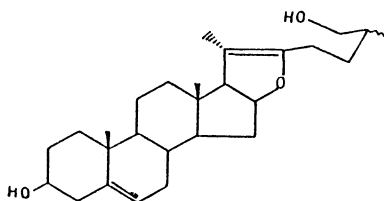
A list of the furostanol saponins reported till 1993, from various plants is tabulated below:



Structure of aglycone (Furostanol)	Corresponding spirostanol (F ring closed)
1. (25S)-5 α -furostan-3 β -amino-22 α , 26-diol	Jurubidine
2. (25S)-5 α -furostan-12-on-3 β , 22 α , 26-triol	Neohecogenin
3. (25R)-5 α -furostan-3 β , 22 α , 26-triol	Tigogenin
4. (25R)-5 α -furostan-2 β , 3 β , 22 α , 26-tetraol	Gitogenin
5. (25R)-5 α -furostan-12-on-3 β , 22 α , 26-triol	Hecogenin
6. (25R)-5 α -furostan-2 α , 3 β , 6 β , 22 α , 26-pentaol	Agigenin
7. (25S)-5 α -furostan-2 α , 3 β , 6 β , 22 α , 26-pentaol	Neoagigenin
8. (25R)-5 α -furostan-3 β , 6 β , 22 α , 26-tetraol	β -Chlorogenin
9. (25R)-5 α -furostan-2 α -OBz, 3 β , 5 α , 6 β , 22 α , 26-pentaol	Karataveginin
10. (25S)-5 α -furostan-3 β , 22 α , 26-triol	Neotigogenin
11. (25R)-5 α -furostan-6-on-3 β , 22 α , 26-triol	Laxogenin
12. (25S)-5 β -furostan-3 β , 22 α , 26-triol	Sarsasapogenin
13. (25R)-5 β -furostan-1 β , 2 β , 3 α , 22 α , 26-pentaol	Tokorogenin
14. (25R)-5 β -furostan-2 β , 3 α , 22 α , 26-tetraol	Yonogenin
15. (25S)-5 β -furostan-2 β , 3 α , 22 α , 26-tetraol	Neoyonogenin
16. (25R)-5 β -furostan-2 β , 3 β , 11 α , 22 α , 26-pentaol	Metagenin
17. (25S)-5 β -furostan-1 β , 3 β , 4 β , 5 β , 22 α , 26-hexaol	Convallagenin-B
18. (25S)-5 β -furostan-1 β , 2 β , 3 β , 4 β , 5 β , 22 α , 26-heptaol	Neopentologenin
19. (25R) -furost-5-en-3 β , 22 β , 26-triol	Diosgenin
20. (25S) -furost-5-en-3 β , 22 α , 26-triol	Yamogenin
21. (25R) -furost-5-en-2 α , 3 β , 22 α , 26-tetraol	Yuccagenin
22. (25R) -furost-5-en-3 β -17 β , 22 α , 26-tetraol	Pennogenin
23. (25R) -furost-5-en-1 β , 3 β , 22 α , 26-tetraol	Rescogenin
24. (25S) -furost-5-en-1 β , 3 β , 22 α , 26-tetraol	Neorescogenin
25. (25R) -furost-5-en-3 β , 17 α , 22 α , 26-tetraol	Gentrogenin
26. (25R) -furost-5-en-3 β , 14 α , 22 α , 26-tetraol	Prazerigenin-A
27. (25S) -furost-5-en-3 β , 14 α , 22 α , 26-tetraol	Neoprazerigenin-A
28. 5 β -furost-25(27)-en-3 β , 22 α , 26-triol	Macranthogenin
29. 5 β -furost-25(27)-en-1 β , 3 β , 22 α , 26-tetraol	Convallamarogenin
30. furost-5, 25(27)-dien-1 β , 3 β , 22 α , 26-tetraol	Δ^5 Convallamarogenin
31. (25R)-5 α -furostan-26-O-acetyl-3 β , 14 α , 17 α , 22 α , 26-pentaol	—
32. (25R)-5 α -furostan-26-O-acetyl-3 β , 17 α , 22 α , 26-pentaol	—
33. (25S)-5 β -furostan-4 β -acetoxo-2 β , 3 α , 22 α , 26-tetraol	Diotigenin-4-acetate



34. 17(20) Dehydrokryptogenin



35. Pseudodiosgenin (25R)

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^{20}$)	Structure
1.	<i>Agave americana</i> (Lf)		
	168, 169	Agavoside-G	$\text{rha-}_3\text{glu-}_4\text{glu-}_4\text{gal-O-}_3[5];$ R = H
	168–170	Agavoside-H 228–30;–113 (MeOH)	$\text{rha-}_4\text{rha-}_3\text{glu-}_4\text{gal-O-}_3[5];$ R = H
2.	<i>Allium ampeloprasus</i>		
	171	Ampeloside-BF ₁	$\text{glu-}_3\text{glu-}_4\text{gal-O-}_3[6];$ R = H
	171	Ampeloside-BF ₂	$\text{glu-}_4\text{gal-O-}_3[6];$ R = H
3.	<i>Allium karataviense</i> (Lf)		
	172	Karatavioside-C	$\text{xyI-}_3\text{glu-}_4\text{gal-O-}_3[21];$ R = H
	173	Karatavegenin- β -glucoside 294–98	$\text{glu-O-}_3[9];$ R = H
4.	<i>Allium narcissiflorum</i>		
	174	Alliumoside-B	$\text{glu-}_3\text{glu-}_3\text{glu-O-}_3[19];$ R = H
	174	Alliumoside-C	$\text{rha-}_4\text{rha-}_4\text{rha-}_6\text{gal-}_6\text{glu-O-}_3[19];$ R = H
	174	Alliumoside-D	$\text{rha-}_4\text{rha-}_6\text{glu-}_3\text{glu-O-}_3[19];$ R = H
	174	Alliumoside-E	$\text{glu-}_4\text{rha-}_4\text{rha-}_6\text{glu-}_3\text{glu-O-}_3[19];$ R = H
	175	Glycoside-B	$\text{glu-}_3\text{glu-}_6\text{glu-O-}_3[19];$ R = H
5.	<i>Allium ostowskianum</i> (Blb)		
	176		$\text{xyI-}_3\text{glu-}_4\text{gal-O-}_3[6, 7];$ R = Me
6.	<i>Allium sativum</i> (Blb)		
	177	Protoeruboside-B	$\text{glu-}_3\text{glu-}_4\text{gal-O-}_3[8];$ R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: [α] _D ⁰)	Structure
7. <i>Allium schubertii</i> (Blb)			
178		Aginoside and Turcoside-A -34.5 (Py)	xyl ³ ₂ glu- ⁴ gal-O- ³ [6, 7]; R = H
8. <i>Allium turcomanicum</i> (Blb)			
179		Turcoside-C	xyl ³ ₂ glu- ⁴ gal-O- ³ [7]; R = H
9. <i>Anemarrhena asphodeloides</i> (Rh)			
180		Anemarsaponin-B 226	glu- ² gal-O- ³ [35];
10. <i>Arecastrum romanzoffianum</i> (Lf)			
181		Methyl proto Pb 192-5; -91.8 (Py)	rha- ⁴ rha ³ ₂ glu- ³ [19]; R = Me
11. <i>Aparagus adscendens</i> (Ft)			
18		Asparoside-A 180-84; -53 (MeOH)	rha ⁴ ₂ glu-O- ³ [12]; R = Me
18		Asparoside-B 170-78; -58 (Py)	-do-; R = H
17		Asparoside-C 167-72; 75 (MeOH)	rha ⁶ ₄ glu-O- ³ [12]; R = Me ara ²
17		Asparoside-D 161-66; -66 (Py)	-do-; R = H
182		Adscendoside-A -89 (MeOH)	rha ⁶ ₄ glu-O- ³ [10]; R = Me rha
182		Adscendoside-B -92 (H ₂ O)	-do-; R = H
12. <i>Asparagus cochinchinensis</i> (Rt)			
121		ASP-IV 165-7; -22.9 (MeOH)	xyl- ⁴ glu-O- ³ [12]; R = Me
121		ASP-IV 146-49; -30.9 (Py)	-do-; R = H
121		ASP-V 150-56; -45.5 (MeOH)	rha- ⁶ glu-O ³ [12]; R = Me
121		ASP-V 270-75; -45.5 (Py)	-do-; R = H
121		ASP-VI 165-68; -50 (MeOH)	rha ⁶ ₄ glu-O ³ [12]; R = Me xyl
121		ASP-VI 169-72; -49.5 (Py)	-do-; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^{20}$)	Structure
121		ASP-VII 179–81; –27 (MeOH)	rha ⁶ xyl ⁴ → ₂ glu–O ³ [12]; R = Me glu
121		ASP-VII 187–90; –31.6 (Py)	—do—; R = H
52		ASP-VIII –76.4 (MeOH)	rha ⁴ glu–O ³ [35];
52		Methylprotodioscin	rha ⁴ rha ² → ₂ glu–O ³ [19]; R = Me
52		Pseudoprotodioscin	rha ⁴ glu–O ³ [19]; R = H
13. <i>Asparagus curillus</i> (Ft)			
25		Curilloside-A 180–84; –53 (MeOH)	rha ⁶ rha ⁴ → ₂ glu–O ³ [12]; R = Me
25		Curilloside-B 170–74; –58 (Py)	—do—; R = H
(Rt) 26		Curilloside-E –47 (MeOH)	ara ⁴ glu ² → ₂ glu–O ³ [12]; R = Me
26		Curilloside-F –51 (H ₂ O)	—do—; R = H
27		Curilloside-C –78	ara ⁶ rha ⁴ → ₂ glu–O ³ [12]; R = Me glu
27		Curilloside-D –66	—do—; R = H
183		Curilloside-G 173–75; –50	glu ⁴ rha ² → ₂ glu–O ³ [12]; R = Me
183		Curilloside-H 181–83; –53	—do—; R = H
14. <i>Asparagus filicinus</i> (Rt)			
184		Aspafiloside-C	ara ⁶ xyl ⁴ → ₂ glu–O ³ [12]; R = Me
184		—	—do—; R = H
15. <i>Asparagus officinalis</i> (Rt)			
77		Officinalisnin-I 162–68; –23.8 (MeOH)	glu ² glu–O ³ [12]; R = H
77		Officinalisnin-II 175–82; –41.6 (MeOH)	xyl ⁴ glu ² → ₂ glu–O ³ [12]; R = H
185		Asparagoside-B	[12]; R = H
186		Asparagoside-E 254–60; –38 (H ₂ O)	glu ³ glu–O ³ [12]; R = H
187		Asparagoside-G	glu ⁴ glu ³ → ₂ glu–O ³ [12]; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^{20}$)	Structure
186		Asparagoside-H	$\text{xyl}-\overset{4}{\text{glu}}-\overset{4}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [12]; R = H
(Sh) 188		Furastanol Saponin-1 192-95; -92.5 (CHCl ₃ : MeOH)	$\text{rha}-\overset{4}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [20]; R = H
188		Furastanol Saponin-2 188-92; -74.6 (CHCl ₃ : MeOH)	$\text{rha}-\overset{4}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [20]; R = H
16. <i>Asparagus plumosus</i>			
189		Furostanol glycoside-I 171-3; -78 (MeOH)	$\text{rha}-\overset{3}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [20]; R = Me
189		Furostanol glycoside-II 183-5; -81 (H ₂ O)	—do—; R = H
17. <i>Asparagus racemosus</i> (Rt)			
190		Asparacemoside-A 213-14	$\text{rha}-\overset{6}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [12]; R = H
190		Asparacemoside-B 180-83	—do—; R = Me
18. <i>Asparagus sprengeri</i> (Rt)			
19		Sprengeroside-A 180-84; -100.3 (MeOH)	$\text{xyl}-\overset{4}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [20]; R = Me
19		Sprengeroside-B 187-92; -99.5 (MeOH)	$\text{rha}-\overset{6}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [20]; R = Me
19		Sprengeroside-C 186-90; -84.5 (Py)	$\text{xyl}-\overset{4}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [20]; R = H
19		Sprengeroside-D 192-98; -83.4 (Py)	$\text{rha}-\overset{6}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [20]; R = H
191		Sprengeroside-I 186-90; -95.5 (Py)	$\text{rha}-\overset{4}{\text{glu}}-\text{O}-\overset{3}{\text{glu}}$ [20]; R = Me
191		Sprengeroside-J 195-98; -79.5 (Py)	—do—; R = H
19. <i>Aspidistra elatior</i>			
192		— 235-6; -34.8 (MeOH)	$\text{glu}-\text{O}-\overset{5}{\text{glu}}$ [17]; R = Me
192		— 195-8; -60.4	$\text{glu}-\text{O}-\overset{5}{\text{glu}}$ [17]; R = H
120		Methylprotoaspidistrin 202-7; -63.4 (Py)	$\text{xyl}-\overset{3}{\text{glu}}-\overset{4}{\text{gal}}-\text{O}-\overset{3}{\text{glu}}$ [19]; R = Me
120		Protoaspidistrin -64 (Py)	—do—; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^{20}$)	Structure
20.	<i>Balanites aegyptica</i> (Ft)		
	193	Balanitoside-A	rha- ² glu- ⁴ glu- ³ [19]; R = Me
	193	Balanitoside-B	—do—; R = H
21.	<i>Balanites roxburghii</i> (Sbk)		
	194	Protodeltonin	glu- ⁴ _{rha} > ² glu-O- ³ [19]; R = H
22.	<i>Beshorneria yuccoides</i> (Lf)		
	195	Beshornoside 219–21; –25 (Py)	rha- ⁴ glu- ³ _{rha} > ² glu-O- ³ [3]; R = H
23.	<i>Capsicum annum</i> (Sd)		
	196	Capsicoside 295; –35 (CHCl ₃ : MeOH)	glu- ³ _{glu} > ² glu- ⁴ gal- ³ glu-O- ³ [4]; R = H
24.	<i>Chamaerops humilites</i> (Wp)		
	197	Methylprotodioscin [Cst ₁ , Cu ₃] 185–89; –102.9 (Py)	rha- ⁴ _{rha} > ² glu-O- ³ [19]; R = Me
	197	Methylproto Pb. [Cl ₂ , Cst ₂ , Cu ₄] 189–90; –86.4 (Py)	rha- ⁴ _{rha} > ² glu-O- ³ [19]; R = Me
	197	Methylprotorhapissaponin [Cu ₅] 196–99; –92.4 (Py)	glu- ⁴ _{rha} - ⁴ _{rha} > ² glu-O- ³ [19]; R = Me
25.	<i>Chlorophytum malayense</i> (Rh)		
	137	Chloromaloside-B 295–98; –48.3 (Py)	xyl- ³ _{glu} > ² glu- ⁴ gal-O- ³ [2]; R = Me
26.	<i>Convallaria majalis</i> (Rt)		
	198	Convallamaroside –66 (MeOH)	rha- ² qui-O- ³ _{glu} > ¹ [29]; R = H
27.	<i>Costus speciosus</i> (Sd)		
	199	Costusoside-I 220–24; –76.5 (Py)	rha- ⁴ _{glu} > ² glu-O- ³ [19]; R = Me
	199	Costusoside-J 248–50; –79.1 (Py)	—do—; R = H
	200	— 194–7; –70.4 (Py)	rha- ² glu-O- ³ [19]; R = Me
	200	Methyl protodioscin 185–7; –102.9 (Py)	rha- ⁴ _{rha} > ² glu-O- ³ [19]; R = Me
	200	Protodioscin 190–6; –57.8 (Py)	—do—; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^{20}$)	Structure
28. <i>Digitalis lanata</i> (Lf)			
32		Lanagitoside 260–83; –18.6 (CHCl ₃ : MeOH)	$\text{glu-}^3_{\text{gal}} \xrightarrow{\text{xy}^1_3} \text{glu-}^4_{\text{gal}} \text{-O-}^3[4];$ R = Me
32		—	—do—; R = H
32		Lanatigotside 268–92; –28.4 (CHCl ₃ : MeOH)	$\text{glu-}^4_{\text{gal}} \xrightarrow{\text{xy}^1_3} \text{gal-O-}^3[3];$ R = H
32		—	—do—; R = H
29. <i>Digitalis purpurea</i> (Lf)			
201		Purpureogitotside	$\text{xy}^1_3 \xrightarrow{\text{glu}} \text{glu-}^4_{\text{gal}} \text{-O-}^3[4];$ R = H
30. <i>Dioscorea collettii</i> (Rh)			
202		Saponin-C 183–5; –81 (H ₂ O)	$\text{rha}^3 \xrightarrow{\text{glu}} \text{glu-O-}^3[20];$ R = H
31. <i>Dioscorea deltoidea</i> (Rh)			
203		Deltoside-II	$\text{glu}^3 \xrightarrow{\text{rha}} \text{glu-O-}^3[19];$ R = H
203		Deltoside-III	—do—; R = Me
32. <i>Dioscorea floribunda</i> (Rh)			
35		Methylprotodioscin	$\text{rha}^4 \xrightarrow{\text{rha}} \text{glu-O-}^3[19];$ R = Me
35		— 182	[19]; R = Me
204		Floribundasaponin-E 226–29; –66 (Py)	$\text{rha-}^3 \text{rha-}^3 \text{rha-}^3 \text{rha-}^4 \text{glu-O-}^3[19];$ R = Me
204		Floribundasaponin-F (243–47); –74.6 (Py)	—do—; R = H
33. <i>Dioscorea gracillima</i> (Rh)			
33		Methylprotodioscin	$\text{rha}^4 \xrightarrow{\text{rha}} \text{glu-O-}^3[19];$ R = Me
33		Protodioscin 190–96; –57.8 (Py)	—do—; R = H
34. <i>Dioscorea tenuipes</i> (Ap)			
205		Diotigenin-4-acetate furanol-26-O-glucoside 172–74; +134.4	[33]; R = Me
35. <i>Dioscorea tokoro</i> (Tc)			
206		Prototokorin 177–78; –31.8 (MeOH)	$\text{glu-O-}^1[12];$ R = H
207		—	[14]; R = H
207		—	[15]; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C; $[\alpha]_D^{20}$)	Structure
36. <i>Dioscorea septemloba</i>			
33		Methylprotogracillin 249–51; –76.9 (MeOH)	glu ³ _{rha} >glu–O ^{–3} [19]; R = Me
33		Protogracillin 235–38; –57.8 (Py)	—do—; R = H
33		Methylprotodioscin	rha ⁴ _{rha} >glu–O ^{–3} [19]; R = Me
33		Protodioscin	—do—; R = H
37. <i>Dioscorea zingiberensis</i>			
208		Zingiberenin-C	rha ³ _{glu} >glu–O ^{–3} [19]; R = H
208		Zingiberenin-D	glu ³ _{rha} >glu–O ^{–3} [20]; R = Me
38. <i>Dracaena afromontana</i> (Twg)			
209		Afromontoside > 300; –69.3 (Py)	rha ⁴ glu–O ^{–3} [19]; R = H But 26–O–rha.
39. <i>Funkia ovata</i> (Lf)			
210		Funkioside-B 258–66; –135 (MeOH)	glu–O ^{–3} [19]; R = H
210–211		Funkioside-I	rha ⁴ _{rha} > _{rha} ⁴ glu ³ glu ⁴ gal–O ^{–3} [19]; xyl R = H
40. <i>Helleborus macranthus</i> (Rt + Rh)			
212		Macranthoside-I 228–31	glu ⁶ glu–O ^{–3} [28]; R = H
212		Macranthoside-I 263–66	—do—; R = Me
41. <i>Heloniopsis orientalis</i> (Wp)			
213		Hb (Methylprotodioscin) 189–92; –98.2	rha ⁴ _{rha} >glu–O ^{–3} [19]; R = Me
213		Hd(III) –92.1	rha ⁴ _{rha} > _{rha} ⁴ glu–O ^{–3} [22]; R = H
213		Hd(III) –100.2	—do—; R = Me
213		Hc(VI)	rha ⁴ _{rha} > _{rha} ⁴ glu–O ^{–3} [34];
42. <i>Kallstroemia pubescens</i> (Wp)			
214		Kallstroemin-A 235–38; –82 (Py)	rha ² rha ² rha ⁶ glu–O ^{–3} [19]; R = H
214		Kallstroemin-B 232–35; –79.5 (Py)	—do—; R = Me
43. <i>Licuala spinosa</i> (Lf)			
215		Methylprotodioscin 166–68; –73 (Py)	rha ³ _{rha} >glu–O ^{–3} [19]; R = Me

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^{20}$)	Structure
44.	<i>Lilium cordatum</i> (Pt)		
	216	Compound-I -47.1 (MeOH)	rha- ² glu-O- ³ [19]; R = Me
	216	Compound-II -75.8 (MeOH)	rha- ² (6-O-acetyl)-glu-O- ³ [19]; R = Me
45.	<i>Lilium hansonii</i> (Blb)		
	217	Compound-9 -48.8 (MeOH)	glu- ⁴ _{rha} > ² glu-O- ³ [3]; R = H
	217	Compound-10	glu- ⁴ _{rha} > ² glu-O- ³ [19]; R = H
46.	<i>Lilium paradarinum</i> (Bib)		
	218-219	Pardarinoside-A -56.4 (MeOH)	rha- ² glu-O- ³ [31]; R = Me But 26-O-Ac.
	218-219	Pardarinoside-B -62 (MeOH)	rha- ² glu-O- ³ [32]; R = Me
	219	Pardarinoside-C -50.5 (MeOH)	glu- ⁴ _{rha} > ² glu-O- ³ [31]; R = Me
	219	Pardarinoside-D -59.7 (MeOH)	glu- ⁴ _{rha} > ² glu-O- ³ [32]; R = Me
	219	Paradarinoside-F -32.8 (MeOH)	ara- ³ _{rha} > ² glu-O- ³ [31]; R = Me
	219	Pardarinoside-G -40.8 (MeOH)	ara- ³ _{rha} > ² glu-O- ³ [32]; R = Me
47.	<i>Lilium regale</i> (Blb)		
	220	—	glu- ³ _{rha} > ² glu-O- ³ [20]; R = Me
48.	<i>Liriope platyphilla</i>		
	221	Methylprotodioscin 185-7; -102.9 (Py)	rha- ⁴ _{rha} > ² glu-O- ³ [19]; R = Me
	221	Protodioscin 190-6; -57.8 (Py)	—do—; R = H
49.	<i>Liriope spicata</i>		
	222	Spicatoside-A	xyl- ³ _{glu} > ² fuc-O- ¹ [23]; R = Me
	222	Spicatoside-B	—do—; R = H
50.	<i>Lycopersicum esculentum</i> (Sd)		
	223	Furastanol saponin 217-20; -24 (MeOH + CHCl ₃)	glu- ² glu- ⁴ gal-O- ³ [10]; R = Me
	223	TFI 217-18; -24 (MeOH + CHCl ₃)	—do—; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: [α] _D ⁰)	Structure
51. <i>Metanartheccium uteoviride</i>			
224	—	166–68; –73 (Py)	ara-O ⁻¹¹ [16]; R = Me
52. <i>Ophiopogon jaburan</i> (Rh)			
225	Glycoside-J ₅	165–68	rha ⁻² (4-O-S-fuc)-O ₃ ^{>1} [23 + 24]; glu-O ₃ ^{>1} [28]; R = H
225	Glycoside-J ₆	201–3	rha ⁻² (4-O-S-ara)-O ₃ ^{>1} [28]; glu-O ₃ ^{>1} [28]; R = H
53. <i>Ophiopogon ohwii</i> (Rh)			
225	Glycoside-O ₆	183–85; –50.8 (Py)	rha ⁻² (4-O-S-fuc)-O ⁻¹ [23]
54. <i>Ophiopogon planiscapus</i>			
226	Glycoside-D	194–7; –70.4 (Py)	rha ⁻² glu-O ⁻³ [19]; R = H
226	Glycoside-F	—	glu ₄ ^{>} rha ₃ glu-O ⁻³ [19]; R = Me
226	—	—	—do—; R = H
226	Glycoside-G	212–5; –55.2 (Py)	rha ⁻² (4-O-S-ara)-O ⁻¹ [23]; R = H
55. <i>Paris formosana</i>			
227	—	174–7; –87	rha ⁻⁴ rha ₄ ^{>} rha ₂ glu-O ⁻³ [19]; R = Me
56.* <i>Paris polyphylla</i> (Tb)			
228	Polyphyllin-G	177–81; –76.9 (Py)	ara ₄ ^{>} rha ₃ glu-O ⁻³ [19]; R = Me
228	Polyphyllin-H	—	—do—; R = H
57. <i>Paris teraphylla</i>			
30, 218	Dehydrokryptogenin tetraglycoside	—	rha ⁻⁴ rha ₄ ^{>} rha ₂ glu-O ⁻³ [34]; R = H
58. <i>Phoenix canariensis</i> (Lf)			
215	Methyl (25S) Proto Pb	255–8; –84.8 (MeOH)	rha ⁻⁴ rha ₄ ^{>} rha ₂ glu-O ⁻³ [20]; R = Me
59.* <i>Phoenix dactylifera</i> (Lf)			
215	Methylprotoprosapogenin-A of dioscin	173–77; –87.6 (Py)	rha ⁻⁴ glu-O ⁻³ [19]; R = Me
215	Methylprotorectinatoside	150–2; –86.7 (Py)	rha ⁻⁵ ara ₄ ^{>} rha ₂ glu-O ⁻³ [19]; R = Me

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^{20}$)	Structure
215		Methylproto Pb 176-9; -83.1 (Py)	$\text{rha-}^4\text{rha}>^4_2\text{glu-O-}^3[19]$; R = Me
60.* <i>Phoenix humilis</i> (Lf)			
215		Methyl (25S) proto Pb 255-58; -84.8 (MeOH)	$\text{rha-}^4\text{rha}>^4_2\text{glu-O-}^3[20]$; R = Me
215		Methyl (25S) proto loureiroside 185-87; -78.3 (Py)	$\text{rha-}^5\text{ara}>^4_2\text{glu-O-}^3[20]$; R = Me
61.* <i>Phoenix loureirii</i> (Lf)			
181		Methylproto Pb 192-95; -91.8 (Py)	$\text{rha-}^4\text{rha}>^4_2\text{glu-O-}^3[19]$; R = Me
181		Methylprotorupicolaside 195-99; -82.6 (Py)	$\text{glu-}^2\text{rha-}^4\text{rha}>^4_2\text{glu-O-}^3[19]$; R = Me
181		Methylprotoloureiroside 183-86; -84.1 (Py)	$\text{rha-}^5\text{ara}>^4_2\text{glu-O-}^3[19]$; R = Me
62.* <i>Phoenix reclinata</i> (Lf)			
181		Methylproto Pb 192-95; -91.8 (Py)	$\text{rha-}^4\text{rha}>^4_2\text{glu-O-}^3[19]$; R = Me
181		Methylprotorupicolaside 195-99; -82.6 (Py)	$\text{glu-}^2\text{rha-}^4\text{rha}>^4_2\text{glu-O-}^3[19]$; R = Me
181		Methylprotoreclinata 165-68; -44.2 (Py)	$\text{rha-}^5\text{ara}>^4_2\text{glu-O-}^3[19]$;
63. <i>Phoenix rupicola</i> (Lf)			
181		Methylprototaccaoside 180-83; -77.2 (Py)	$\text{rha-}^3\text{glu-O-}^3[19]$; R = Me
181		Methylproto Pb 192-95; -91.8 (Py)	$\text{rha-}^4\text{rha}>^4_2\text{glu-O-}^3[19]$; R = Me
181		Methylprotorupicolaside 195-99; -82.6 (Py)	$\text{glu-}^2\text{rha-}^4\text{rha}>^4_2\text{glu-O-}^3[19]$; R = Me
64. <i>Polygonatum kingianum</i> (Rh)			
229		Kingianoside-C -27.4 (Py)	$\text{glu-}^4\text{gal-O-}^3[25]$; R = Me
229		Kingianoside-C -34.4 (Py)	$\text{gal-}^4\text{fuc-O-}^3[25]$; R = H
229		(25R) epimer of PO-8 -59.4 (Py)	$\text{glu-}^4\text{gal-O-}^3[19]$; R = Me
65. <i>Polygonatum latifolium</i> (Lf)			
230		Potopolygonatoside-E	$\text{glu-}^3\text{glu-}^4\text{gal-}^3\text{glu-O-}^3[19]$; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^0$)	Structure
66. <i>Polygonatum odoratum</i>			
231	PO-C	208–10; –49 (Py)	$\text{glu}_3\text{glu}_2\text{glu}_4\text{gal-O}^{-3}$ [27]; R = Me
231	PO-d	208–13; –48.1 (Py)	$\text{glu}_3\text{glu}_2\text{glu}_4\text{gal-O}^{-3}$ [26 + 27]; R = Me
67. <i>Polygonatum prattii</i> (Rt)			
232	Pratioside-B	–52.5 (H ₂ O)	$\text{glu}_2\text{glu}_4\text{gal-O}^{-3}$ [25]; R = H
68. <i>Polygonatum officinale</i> (Rh)			
233	Polyfuroside-β	252–58; –68 (Py)	$\text{glu}_4\text{glu}_2\text{glu}_4\text{gal-O}^{-3}$ [19]; R = H
69. <i>Polygonatum verticillatum</i>			
234	Methyl protodioscin		$\text{rha}_4\text{glu}_2\text{glu-O}^{-3}$ [19]; R = Me
234	Protodioscin		—do—; R = H
70. <i>Radix sarsaparilla</i>			
22, 31	Sarsaparilloside	–44 (H ₂ O)	$\text{rha}_6\text{glu}_4\text{glu-O}^{-3}$ [12]; R = H R = H
71. <i>Rhapis exelsa</i> (Wp)			
235	Methylprotodioscin [Est ₂ , El ₁ , Eu ₂]	185.90; –102.9 (Py)	$\text{rha}_4\text{glu}_2\text{glu-O}^{-3}$ [19]; R = Me
235	Methylprotodeltonin [Est _{4-a}]	196–99; –59.8 (Py)	$\text{glu}_4\text{glu}_2\text{glu-O}^{-3}$ [19]; R = Me
235	Methylproto Pb [El ₂ , Est _{4-b} , Eu ₃]	189–90; –86.4 (Py)	$\text{rha}_4\text{glu}_2\text{glu-O}^{-3}$ [19]; R = Me
235	Methylprotorhapissaponin [El ₃]	196–99; –92.4 (Py)	$\text{glu}_4\text{rha}_4\text{glu}_2\text{glu-O}^{-3}$ [19]; R = Me
72. <i>Rhapis humilis</i> (Wp)			
235	Methylprotoprosapogenin- A of dioscin [Hst ₄ , Hl ₁ , Hu ₂]	178–80; –81.2 (Py)	$\text{rha}_4\text{glu-O}^{-3}$ [19]; R = Me
235	Methylprotodioscin [Hl ₂ , Hst ₅ , Hu ₂]	185–89; 102.9 (Py)	$\text{rha}_4\text{glu}_2\text{glu-O}^{-3}$ [19]; R = Me

S. No.	Source (Part) with reference	Saponin (m.p. °C; $[\alpha]_D^{20}$)	Structure
235		Methylprotodeltonin [Hst ₆] 196–99; –59.8 (Py)	glu ⁴ _{rha} > ₂ glu–O ^{–3} [19]; R = Me
235		Methylproto Pb [Hl ₃ , Hu ₄] 189–90; –86.4 (Py)	rha ^{–4} _{rha} > ₄ glu–O ^{–3} [19]; R = Me
73. <i>Ruscus aculeatus</i> (Rh)			
236		Deglucoscoside	rha ^{–2} ara–O ^{–1} [30]; R = H
236		Ruscocide	glu ^{–3} rha ^{–2} ara–O ^{–1} [30]; R = H
74. <i>Smilax aristolochiaefolia</i> (Rt)			
22, 31		Sarsaparilloside –44 (H ₂ O)	glu ⁶ _{rha} > ₄ glu–O ^{–3} [12]; R = H rha ₂
75. <i>Smilax aspera</i> (Lf)			
237		Asperoside	rha ^{–4} _{rha} > ₄ glu–O ^{–3} [20]; R = H
76. <i>Smilax chinae</i>			
33		Smilax saponin-B 183–86; –90 (MeOH)	rha ^{–4} _{rha} > ₄ glu–O ^{–3} [19]; R = Me
(Rh and Rt) 238		Pseudoprotodioscin –72 (MeOH)	rha ⁴ _{rha} > ₂ glu–O ^{–3} [35];
238		Methyl protodioscin –86 (MeOH)	rha ⁴ _{rha} > ₂ glu–O ^{–3} [19]; R = Me
77. <i>Smilax sieboldi</i> (Rh)			
239		Compound-4 –44.8 (EtOH)	ara ^{–6} glu–O ^{–3} [11]; R = H
239		Compound-5 –46.3 (EtOH)	ara ⁶ glu ⁴ > ₂ glu–O ^{–3} [11]; R = H
78. <i>Solanum lyratum</i> (Ber)			
249		Methylprotoaspidistrin 202–7; –63.4 (Py)	rha ³ _{glu} > ₂ glu–O ^{–3} [19]; R = Me
240		Aspidistrin 210–14; –64 (Py)	—do—; R = H
241		Furastanol glucuroside –61.4 (H ₂ O)	glu ³ _{rha} > ₂ glu–O ^{–3} [19]; R = H
79. <i>Solanum melangena</i> (Sd)			
242		Melongoside-N 187–89; –15 (MeOH)	glu ^{–2} glu–O ^{–3} [3]; R = H
242		Melongoside-O 183.–84; –19 (MeOH)	glu ^{–2} glu–O ^{–3} [19]; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^{20}$)	Structure
	242	Melongoside-P 179–80; –75 (H ₂ O)	rha ³ _{glu} > ₂ glu–O ^{–3} [3]; R = H
80.	<i>Solanum nigrum</i> (Wp)		
	243	Uttroside-A 220–25; –49 (MeOH)	xyl ³ _{glu} > ₂ glu– ⁴ gal–O ^{–3} [3]; R = Me
	243	Uttroside-B 210–15; –46 (Py)	—do—; R = H
81.	<i>Solanum paniculatum</i> (Rt)		
	20–21	Jurubine 212–14; –31 (Py)	[1]; R = H
82.	<i>Trachycarpus fortune</i>		
	244	Methylprotodioscin 185–89; –102.9 (Py)	rha ⁴ _{rha} > ₂ glu–O ^{–3} [19]; R = Me
	244	Protodioscin	—do—; R = H
	244	— 174–7; –87	rha ⁴ _{rha} – ⁴ rha> ₂ glu–O ^{–3} [19]; R = H
83.	<i>Trachycarpus wagnerianus</i> (Wp)		
	197	Methylprotodioscin [Tst ₁] 185–89; –102.9 (Py)	rha ⁴ _{rha} > ₂ glu–O ^{–3} [19]; R = Me
	197	Methylproto Pb [Tl ₃ , Tu ₃] 189–90; –86.4 (Py)	rha ⁴ _{rha} – ⁴ rha> ₂ glu–O ^{–3} [19]; R = Me
	197	Pseudoprotodioscin [Tst ₂] 174–76; –80.4 (Py)	rha ⁴ _{rha} > ₂ glu–O ^{–3} [35]
	197	Pseudoproteo Pb [Tl ₄ , Tu ₄] 181–83; –84.4 (Py)	rha ⁴ _{rha} – ⁴ rha> ₂ glu–O ^{–3} [35]
84.	<i>Tribulus terrestris</i>		
	245–247	Methylprotodioscin 189–90	rha ⁴ _{rha} > ₂ glu–O ^{–3} [19]; R = Me
	245–247	Protodioscin	—do—; R = H
	245–247	Methylprotogracillin	glu ³ _{rha} > ₂ glu–O ^{–3} [19]; R = Me
85.	<i>Trigonella coerulea</i> (Sd)		
	245–247	Methylprotodioscin 189–93	rha ⁴ _{rha} > ₂ glu–O ^{–3} [19]; R = Me
	245–247	Protodioscin	—do—; R = H
86.	<i>Trigonella foenumgraecum</i> (Sd)		
	247–249	Trigonelloside-B	rha ⁴ _{rha} > ₂ glu–O ^{–3} [20]; R = Me
	247–249	Trigonelloside-C	—do—; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: [α] _b)	Structure
250		Furostanol glycoside 242–46	glu ^{>3} _{rha} 2glu-O ⁻³ [10]; R = Me
48		Trigofoenoside-A-1 219–21; -90.1 (Py)	rha ⁻² glu-O ⁻³ [20]; R = Me
48		Trigofoenoside-A 210–13	—do—; R = H
251		Trigofoenoside-B 198–200; -62.1 (Py)	rha ⁻⁴ glu-O ⁻³ [10]; R = Me
251		Trigofoenoside-C 210–12; -64.1 (Py)	rha ^{>4} _{rha} 2glu-O ⁻³ [4]; R = Me
48		Trigofoenoside-D-1 246–48; -73.2 (Py)	glu ^{>3} _{rha} 2glu-O ⁻³ [20]; R = Me
48		Trigofoenoside-D	—do—; R = H
252		Trigofoenoside-F-1 256–58; -78.9 (Py)	rha ⁻² glu ⁻⁶ glu-O ⁻³ [19]; R = Me
252		Trigofoenoside-F 233–36	—do—; R = H
252		Trigofoenoside-G-1 270–74; 79.2 (Py)	xyl ^{>4} _{rha} 2glu ⁻⁶ glu-O ⁻³ [19]; R = Me
252		Trigofoenoside-G 275–8	—do—; R = H
253		—	xyl ^{>4} _{rha} 2glu-O ⁻³ [3]; R = Me
87.	<i>Trillium kamschaticum</i> (Ugp)		
30		Furastanol bisglycoside 265–71; -83.1 (Py)	rha ⁻² glu-O ⁻³ [19]; R = Me
30		Protopennogenin, 3-O- rhamnosyl glucoside 275–80; -84.6 (Py)	rha ⁻² glu-O ⁻³ [22]; R = H
30		Dehydrokryptogenin- diglycoside 265–68; -80.1 (Py)	rha ⁻² glu-O ⁻³ [34]; R = H
254		Tj 205–12; -66.5	rha ^{>4} _{rha} 2glu-O ⁻³ [22]; R = Me
254		Tk 194–200; -90 (Py)	rha ⁻⁴ rha ^{>4} _{rha} 2glu-O ⁻³ [22]; R = Me
88.	<i>Trillium tschonoskii</i> (Rt)		
255		Methylprotodioscin	rha ^{>4} _{rha} 2glu-O ⁻³ [19]; R = Me
89.	<i>Yucca filamentosa</i> (Rt)		
34		Protoyuccoside-C 182–84; -30 (MeOH)	gal ⁻² glu ⁻⁴ glu-O ⁻³ [12]; R = H
256		Protoyuccoside-E 150–52; -29 (MeOH)	gal ^{>6} _{gal} 2glu ⁻⁴ glu-O ⁻³ [12]; R = H

S. No.	Source (Part) with reference	Saponin (m.p. °C: $[\alpha]_D^0$)	Structure
90.	<i>Yucca gloriosa</i> (Fl) 257	YG-4 + 21 (Py)	$\text{xyl}^{-3}\text{glu} >_2^{\text{xyl}} \text{glu}^{-4}\text{gal}-\text{O}^{-3}[4];$ R = H

The following abbreviations have been used:

1. glu = D-Glucose; ara = L-Arabinose; xyl = D-Xylose; gal = D-Galactose; rha = L-Rhamnose, qui = D-Quinovose; fuc = Fucose; Rt = Root; Rh = Rhizome; Lf = Leaf; Sd = Seed; Ugp = Underground part; Wp = Whole part; Blb = Bulb; Ap = Aerial part; Ft = Fruit; Sh = Shoot; Sbk = Stem bark; Tc = Tissue culture; Twg = Twigs; Tb = Tuber; Ber = Berry; Tl = Flower; Pt = petals.
- 2.* All the sugars reported in the above saponins appear in the pyranose form except* marked, where the sugars reported are in the furanose form.
3. The number in the brackets refer to the structure of aglycone part.
4. The linkages of sugars with each other, and with aglycone are β .

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