



Photocatalytic Degradation and Anticancer Activity of Green Synthesized Molybdenum Nanoparticles for Inhibiting the Cervical Cancer Cells

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This work reports the synthesis of molybdenum nanoparticles (Mo NPs) through green routes using horse gram seed extracts and characterized by GC-MS, electronic (UV-visible) spectroscopy, IR, XRD and SEM techniques. Their potent applications as *in vitro* anticancer agent against HeLa cell lines and photocatalytic degradation of methyl orange (MO) and methyl violet (MV) dyes were also evaluated. The anti-malignant properties of Schiff base ligands [2-fluorobenzaldehyde semicarbazone (L₁H) and 2-fluorobenzaldehyde thiosemicarbazone (L₂H)] with molybdenum metal precursor [dioxobis(2,4-pentanedionato)molybdenum] were also explored and their Mo complexes after conversion into nano form as both ligands, L₁H and L₂H approve their binding property based on molecular docking studies.

Keywords: Green synthesis, Molybdenum nanoparticles, Antioxidant activity, Schiff base, Metal complexes, Anticancer activity.

INTRODUCTION

Nanotechnology has emerged as a powerful tool in many fields of science and engineering due to its focus on altering and enhancing materials on a nanoscale scale [1]. It is a fascinating domain to alter the materials in size, shape, morphology and dimensions, creating nanoparticles almost equal to the Bohr radius [2]. Green technology, which includes the employment of microorganisms like bacteria, fungi, yeast and plant material extract as a safe and eco-friendly tool for synthesizing nanoparticles, has taken the lead over physical and chemical methods [3,4]. Therefore, biogenic preparation of metal nanoparticles is extremely popular nowadays. Owing of existence of phytochemicals like flavonoids, terpenoids and quinines phenols, alkaloids and tannins plant-mediated synthesis has become a powerful unconventional instrument for synthesizing nanoparticles [5,6]. The phytonutrients are playing significant role in governing particle size *via* prompting and topographical reactions and performing role of capping agents [7].

Molybdenum has gained much attention owing to its wide-range applications [8,9]. Since molybdenum exist in various oxidation states and having low toxicity than other *d*-block elements thus, molybdenum nanoparticles are made to absorb,

to cure many deficiency disorders. The deficiency may sometimes disrupt enzyme activity in nitrogen metabolism [10,11].

Molybdenum oxide derivatives due to their distinctive electrochemical and diverse topography is one of the captivating transition metal oxides [12]. Molybdenum oxide a verified chemical for various uses, primarily for Li-ion battery [13]. Still, a few reports have also used them to decontaminate water [14]. Even though several reports are available which are explaining the amazing photocatalytic activity of molybdenum trioxide (MoO₃) [15], though, few studies are there to examine the photocatalytic activity of MoO₂ [16]. Surface active ingredients are important tools to carve the morphological structure of nanoparticles, which in turn alters the property of material [17]. A little research support is there to describe the geomorphological controlled preparation of molybdenum oxide [18].

Of late, MoO₃ nanostructures display photocatalytic properties under UV-visible light. With the nanoparticles of MoO₃ under visible light, several applications have been explored and examined, such as photosensitive deterioration of methylene violet dye [19,20] and methylene orange dye [21], gas sensing properties [22], supercapacitor [23], oxidation of methanol [24], epoxidation [25], light/optical properties [26], lithium

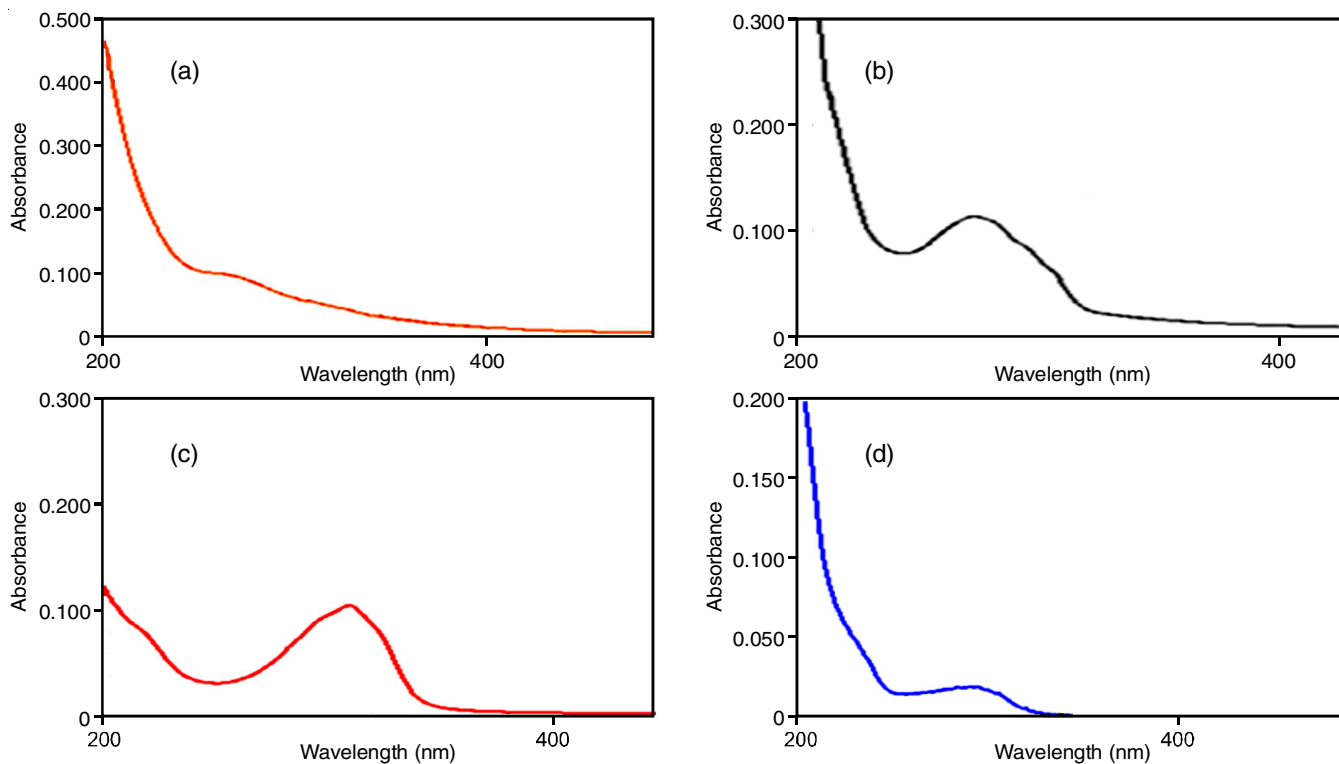


Fig. 2. Electronic spectra of (a) L₁H, (b) L₂H, (c) ML₁, (d) ML₂

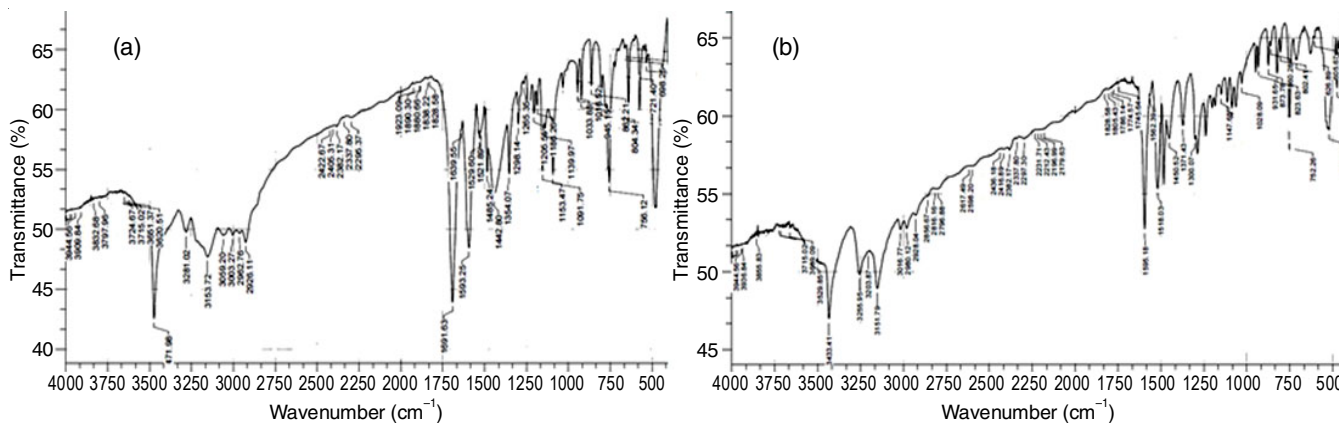


Fig. 3. IR spectra of NPs (a) L₁H (b) L₂H

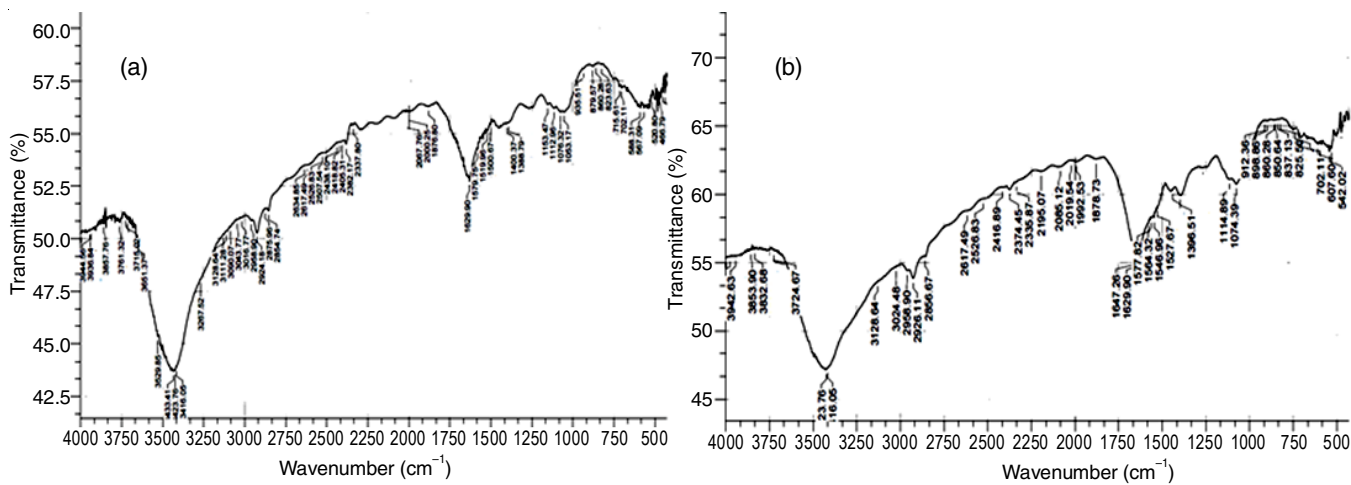


Fig. 4. IR spectra of NPs (a) ML₁ (b) ML₂

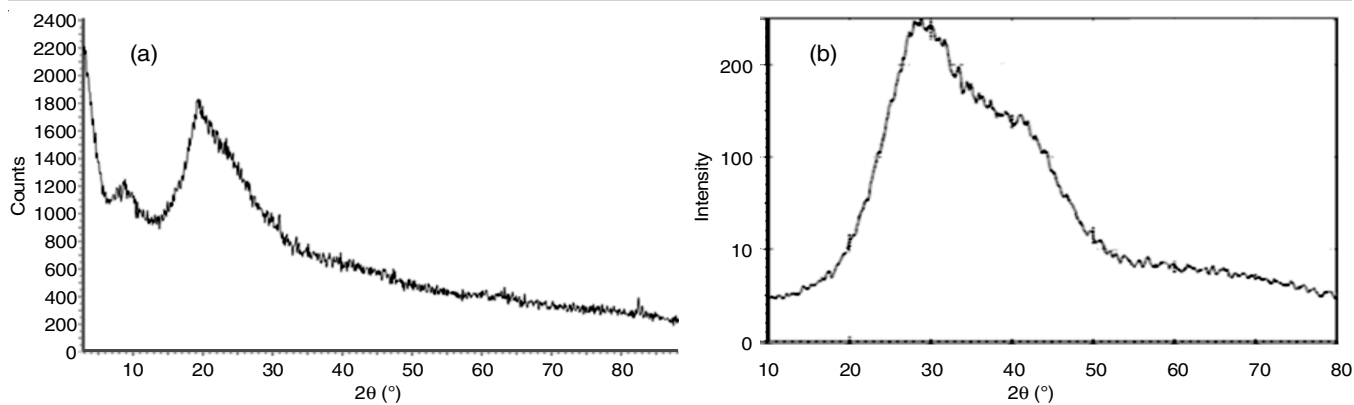


Fig. 5. XRD spectra of (a) Mo-NPs ML₁ and (b) Mo-NPs ML₂

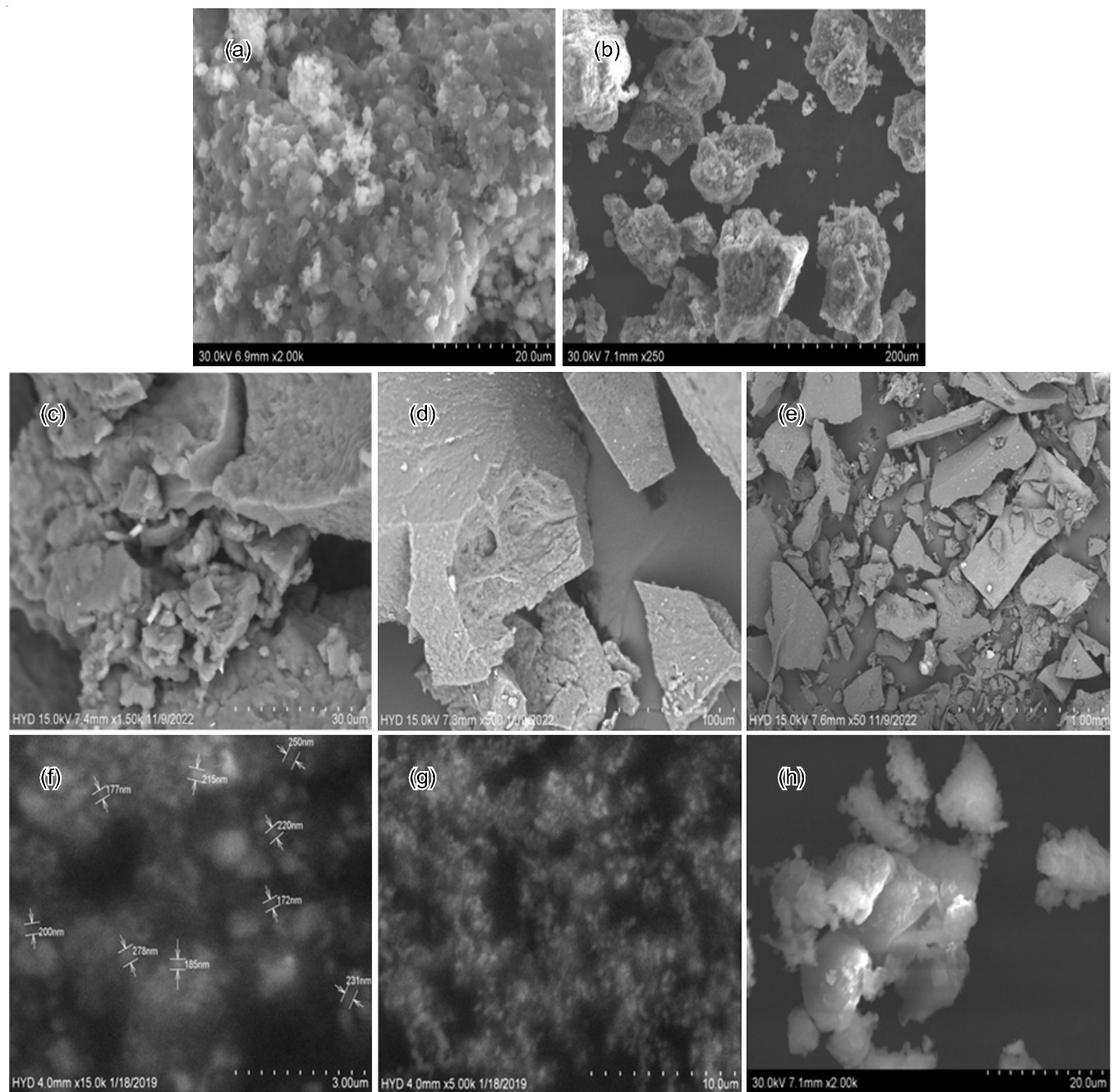


Fig. 6. SEM images of (a) L₁H, (b) L₂H, (c-e) ML₁ and (f-h) ML₂

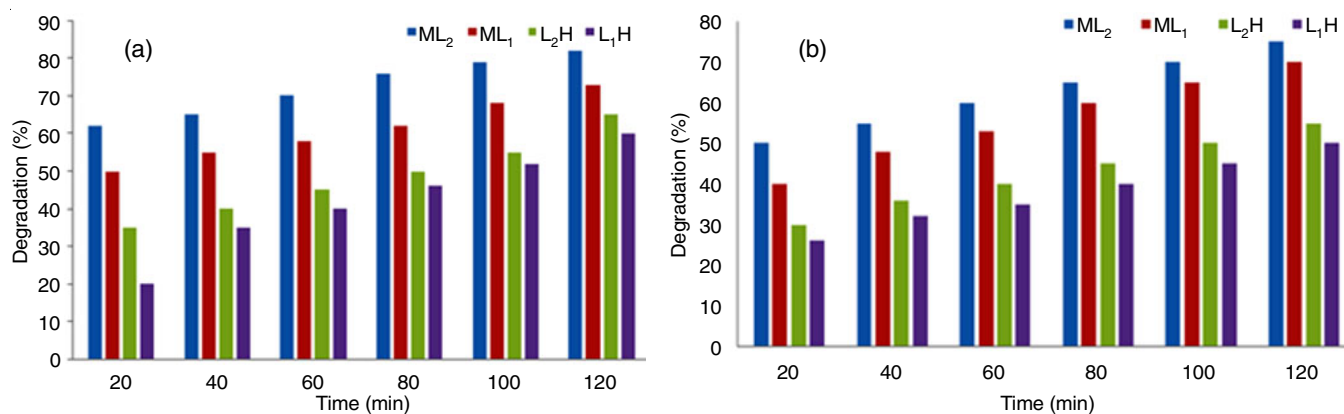


Fig. 7. Bar graphs showing photocatalytic activity of all NPs with (a) methyl orange and (b) methyl violet dyes

Molecular docking studies: Molecular docking study were performed using LibDock to study analyze the selected complex binding approach and affinity with modeled RAT nitric oxide synthase of brain protein. Based on the LibDock score and extremely interrelating amino acid residues, the finest ligand conformation was selected. Rutin exhibits the highest LibDock score of 119.72 Kcal/mol and binding affinity among the 10 conformations generated for each complex. The interaction analysis of H-bonding affinity with nitro-L-arginine methyl ester shows 90.235 Kcal/mol. Apart from rutin and other compounds, hesperidin also shows good binding affinity to LibDock score count 112.331 Kcal/mol (Table-2). In this work, we have analyzed compounds based on selected target proteins for their anti-malignant activity through molecular

docking study [40]. The binding interaction of complexes L_1H and L_2H with the target protein was confirmed using docking results, indicating significant interactions and implying their inhibitory effect (Fig. 8). This study suggests that compounds L_1H and L_2H could potentially inhibit RAT nitric oxide synthase in brain proteins related to growth and may serve as a therapeutic drug.

Anticancer activity: Evaluation of cell viability for green synthesized Mo NPs was performed using MTT assay against HeLa cancer cell lines. The cell viability of the cancer cell depends on the dosage of the compounds. The results were compared with the standard cisplatin; the IC_{50} value of cisplatin against the HeLa cell line was found to be 3.58 $\mu\text{g}/\text{mL}$. Among the four complexes, ML_2 complex exhibits the most significant

TABLE-2
DOCKING SCORES OF THE COMPOUNDS WITH MODELED RAT NITRIC OXIDE SYNTHASE OF BRAIN PROTEIN

Name of the molecule	C docker energy	-c docker interaction energy	Interacting amino acids	Interacting atoms	H-Distance
Ligand 1 L_1H	30.259	32.722	DG12, DT10, Asn722, Thr718, TGp11, Asp533, Arg364, DA113, Dc112	A: ARG364: HH22 - Ligand1: O13	1.831000
Ligand 2 L_2H	48.512	58.769	DG12, DT10, Asn722, Thr718, TGp11, Asp533, Arg364, DA113, Dc112	A: THR718: HG1 - Ligand2: F7 Ligand2: H20 - A: ASP533: OD2 Ligand2: H19 - A: ASP533: OD2	2.307000 2.468000 2.222000

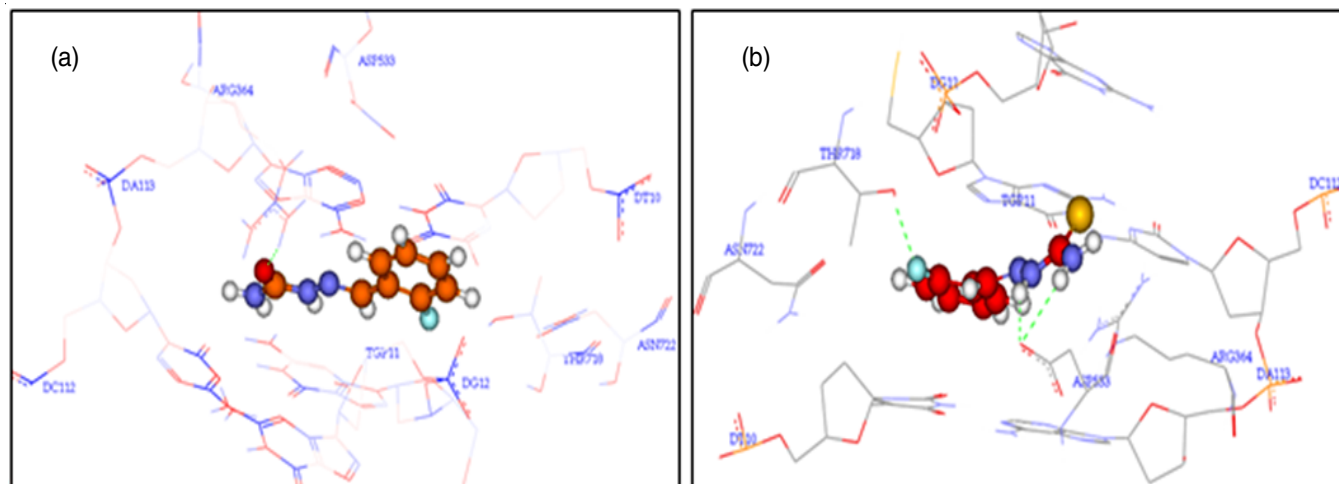


Fig. 8. H-bond interactions of (a) L_1H and (b) L_2H . H-bond represented by green dotted lines and letters showing the amino acids involved in the bonding and compounds are shown by the ball model

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