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Molecular docking analysis of docetaxel analogues as dual lipocalin 2 inhibitors

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Abstract

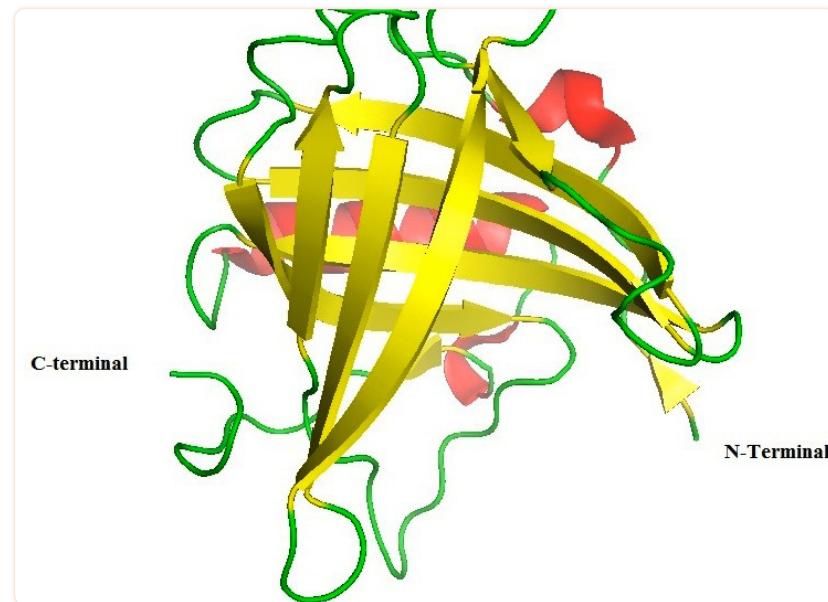
Lipocalin 2 (Lcn2, also called as neutrophil gelatinase-associated lipocalin) is a member of the lipocalin family and a known target for breast cancer. Therefore, it is of interest to use Docetaxel as a scaffold to design molecules with improved efficiency from naturally derived phytochemicals. We document 10 analogues (4Deacetyltaxol, 7Acetyltaxol, Cabazitaxel, Cephalomannine, Docetaxal, Deacetyltaxol, Docetaxeltrihydrate, Ortataxel, Paclitaxel, Taxoline) having optimal binding with Lipocalin 2 in comparison with Docetaxel. This data is highly useful for consideration in the design and development of drugs for breast cancer.

Keywords: Lipocalin 2, docetaxel, analogues, molecular docking

Background

Breast cancer is an issue of medical importance worldwide [1-3]. Treatments such as radiation therapy, chemotherapy, surgery, immunotherapy, and hormone therapy are available with debatable efficiency. Known drugs in this context is under constant debate for efficiency and drug resistance [4,5]. The use of an FDA approved drug docetaxel as a therapeutic agent in cancer patients are known [6-10]. Lipocalin 2 (Lcn2, neutrophil gelatinase-associated lipocalin ([Figure 1](#)) is a member of the lipocalin family and a known target for breast cancer [11-18]. Therefore, it is of interest to use Docetaxel as a scaffold to design molecules with improved efficiency from naturally derived phytochemicals.

Feedback



[Figure 1](#)

Structure of lipocalin 2

Methods:

Protein preparation:

The X-ray crystallographic structure of the lipocalin 2 with 2.6 Å resolution was retrieved from Protein Data Bank (PDB) with PDB ID: 1DFV was used in this study using standard procedure [\[19\]](#).

Ligand preparation:

Structure of Docetaxel and its 10 analogues were downloaded from the PUBCHEM database in SDF format and converted to PDF file format with the help of the Online Smile Translator.

Molecular docking analysis:

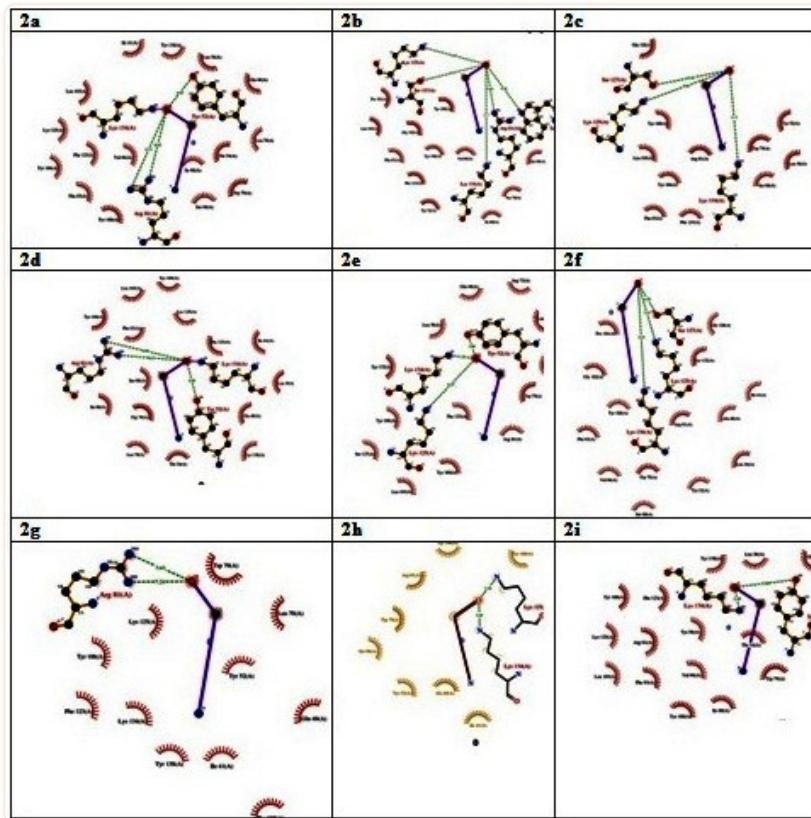
Molecular docking analysis was completed using PATCHDOCK following standard protocols [\[20,21\]](#). The docked structure was examined using Ligplot [\[22\]](#).

Results and discussion:

[Table 1](#) shows the Molecular docking analysis of Docetaxel analogues as dual Lipocalin 2 inhibitors. We document 10 analogues (4Deacetyltaxol, 7Acetyltaxol, Cabazitaxel, Cephalomannine, Docetaxal, Deacetyltaxol, Docetaxeltrihydrate, Orataxel, Paclitaxel, Taxoline) with desirable binding with the Lipocalin 2 in comparison with Docetaxel ([Table 1](#)). Results of the analogue deacetyltaxol have the good binding energy (-132-89 kcal/mol). [Figure 2](#) shows ligand-protein interaction drawn using LigPlot. The interacting residues with optimal hydrogen bonding patterns are shown. An increased amount of hydrophobic atoms in the active center of drug-target boundary enlarged the biological action of the lead [\[23\]](#).

Table 1**Molecular docking analysis of docetaxel analogues as dual lipocalin 2 inhibitors**

| S. No | Compound name | Score | ACE | Atomic interaction | Ligand atom | Distance | No of non bonded interaction |
|-------|---------------------|-------|---------|---|---------------------------------------|-------------------------------------|------------------------------|
| 1 | Docetaxel | 5804 | -54.82 | LYS 125 LYS 134 | NZ-O | 1.53 3.02 | 57 |
| 1 | 4Deacetyltaxol | 6474 | -147.98 | TYR 52 ARG 81 LYS 134 | OH-O NH-0 NZ-O | 2.87 1.49 3.32 | 117 |
| 2 | 7Acetyltaxol | 6252 | -103.92 | TRP 79 ARG 81 LYS 125 SER 127 LYS 134 | NE-O NH2-O NZ-O OG-O NZ-O | 2.3 3.29 2.83 1.39 3.14 | 114 |
| 3 | Cabazitaxel | 5952 | -50.11 | LYS 125 SER 127 LYS 134 LYS 34 | NZ-O OG-O NZ-O NZ-O | 2.24 2.44 3.29 3.03 | 69 |
| 4 | Cephalomannine | 6794 | -113.10 | TYR 52 ARG 81 ARG 81 LYS 134 | OH-O NH1-O NH2-O NZ-O | 2.62 1.43 2.17 1.84 | 110 |
| 5 | Docetaxal | 6404 | -111.73 | TYR 52 TYR 52 LYS 125 LYS 134 | OH-O OH-O NZ-O NZ-O | 3 2.81 3.28 2.63 | 108 |
| 6 | Deacetyltaxol | 5694 | -132.89 | LYS 125 SER 127 LYS 134 | NZ-O OG-O NZ-O | 3.04 2.68 2.05 | 87 |
| 7 | Docetaxeltrihydrate | 6022 | -63.23 | ARG 81 ARG 81 | NH1-O NH2-O | 2.39 1.34 | 84 |
| 8 | Ortataxel | 6204 | -55.51 | LYS 125 | NZ-O | 2.26 | 74 |

**Figure 2**

Ligplot analysis of docked complex showing interaction of lipocalin 2 with (a) 4Deacetyltaxol; (b) 7Acetyltaxol; (c) cabazitaxel; (d) Cephalomannine; (e) Docetaxal; (f) Deacetyltaxol; (g) Docetaxeltrihydrate; (h) ortataxel; (i) paclitaxel; (j) taxoline

Conclusions:

We document 10 analogues (4-deacetyltaxol, 7-acetyltaxol, cabazitaxel, cephalomannine, docetaxal, deacetyltaxol, docetaxeltrihydrate, ortataxel, paclitaxel and taxoline) with desirable binding features with the Lipocalin 2 in comparison with Docetaxel for further in vivo and in vitro validation.

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