

# **siRNAmod: A database of experimentally validated chemically modified siRNAs**

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## **Additional files**

### **Additional file 1**

**Figure S1.** Distribution of natural and non-natural chemical modifications.

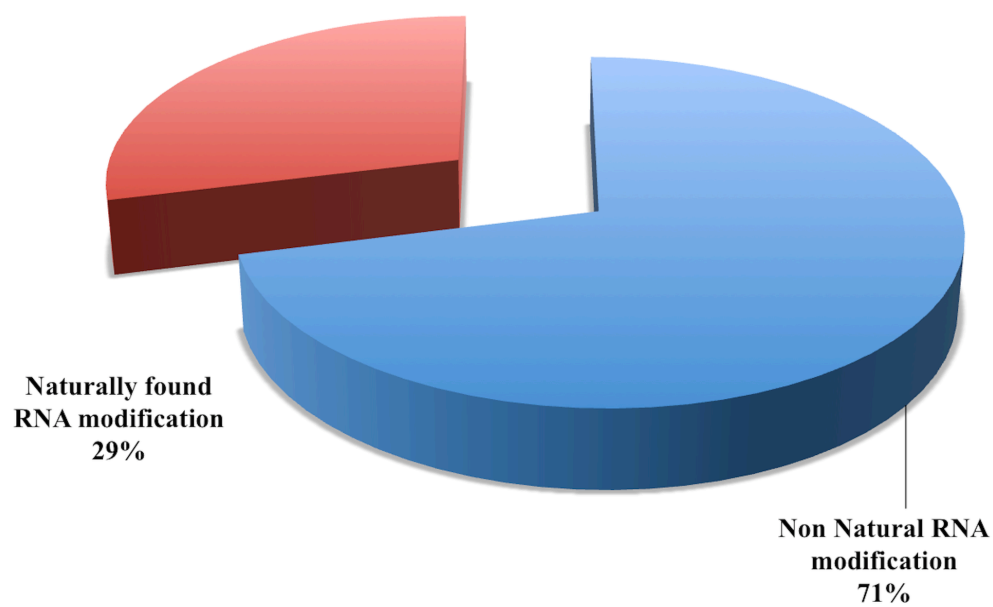
**Figure S2.** Relative frequency of chemical modifications on sense and antisense cm-siRNAs for 21 base pair.

**Figure S3.** Cumulative frequency of cm-siRNA modifications on sense and antisense sequences.

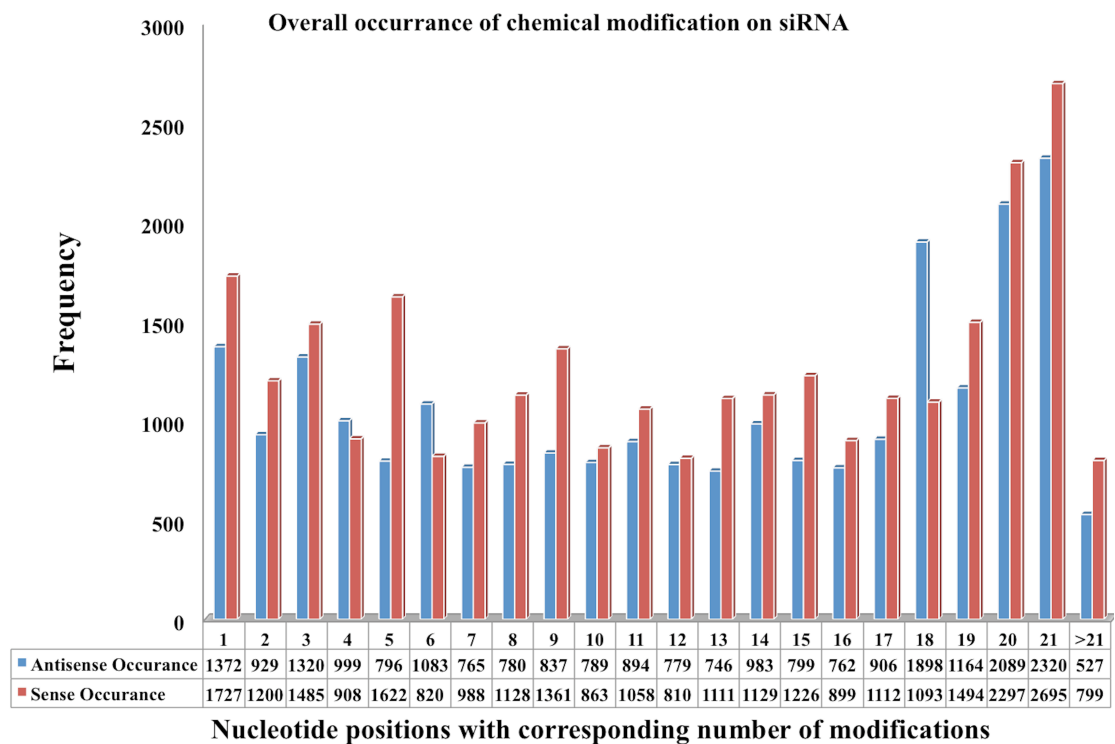
**Figure S4.** Grouping frequency of different unique chemical modifications.

**Figure S5.** Cell lines distribution in siRNAmdb database.

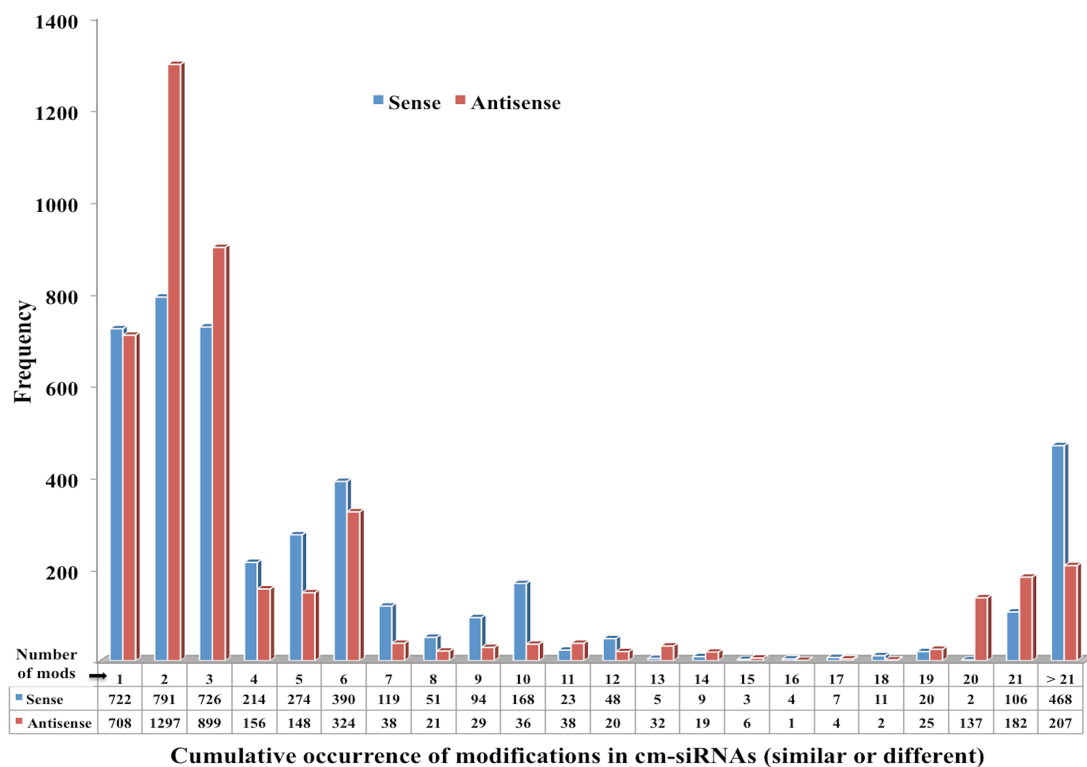
### Natural and non natural based distribution of chemical modifications



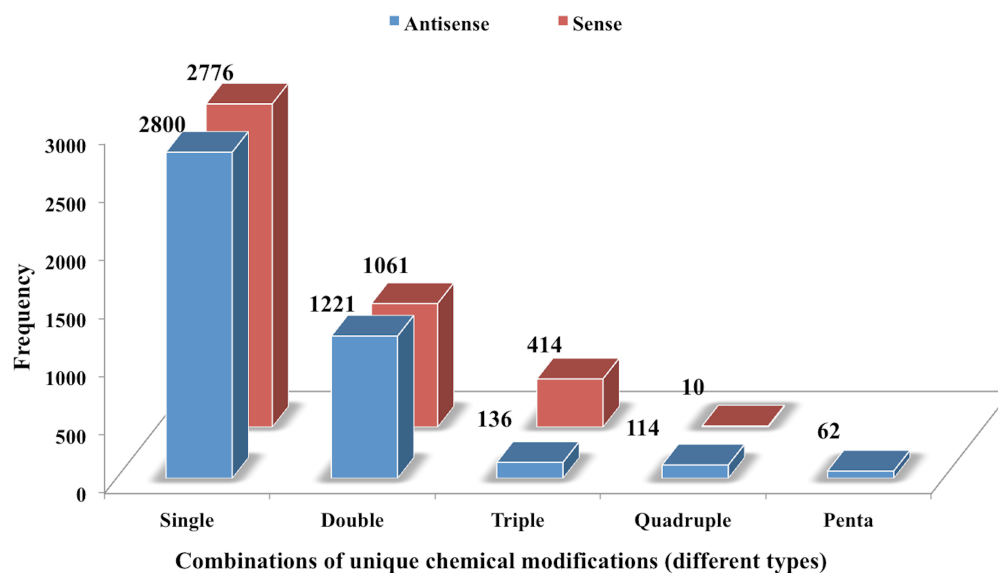
**Figure S1.** Distribution of natural and non-natural chemical modifications.



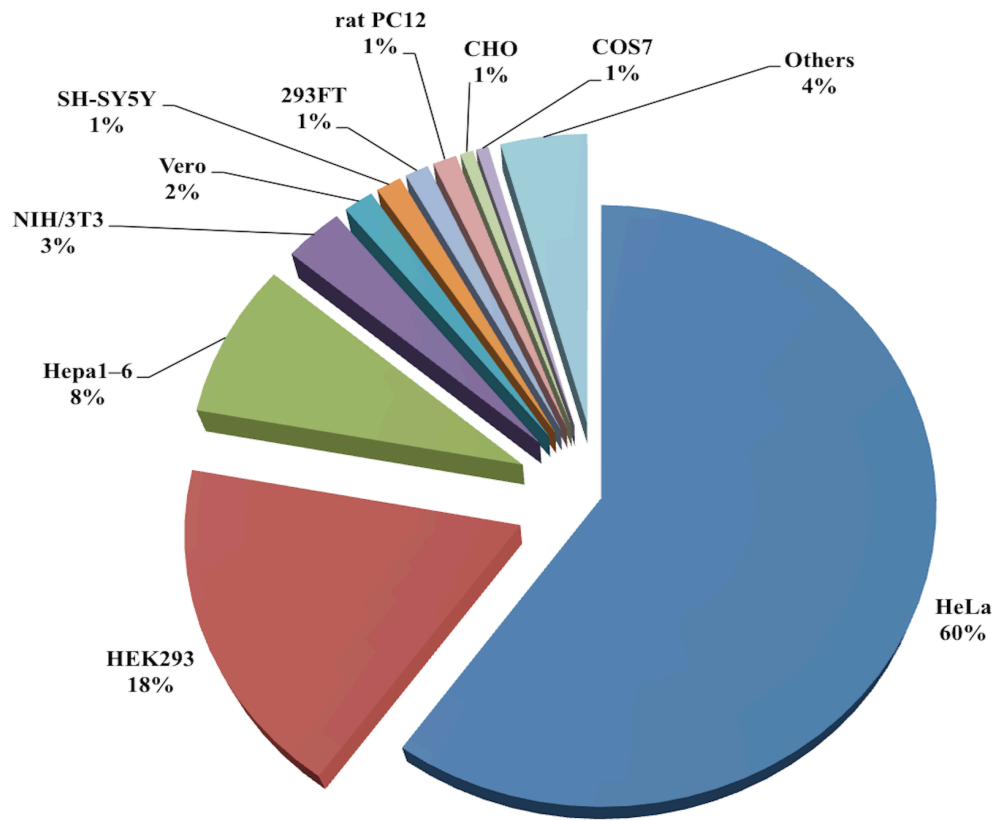
**Figure S2.** Relative frequency of chemical modifications on sense and antisense cm-siRNAs for 21 base pair. Nucleotide position represented as 1 to 21 both starting from 5' end on X-axis. >21 signifies end modifications or additional modifications attached after 21<sup>st</sup> nucleotide. Further number of each position is provided below its corresponding position.



**Figure S3.** Cumulative frequency of cm-siRNA modifications on sense and antisense sequences. Number of Mods on X-axis represents the cumulative number of modifications on cm-siRNAs i.e. total number of modifications (same or different) are present on the particular cm-siRNAs. Their respective numbers are also provided in proceeding rows on X-axis.



**Figure S4.** Grouping frequency of different unique chemical modifications i.e. how many different modifications are grouped in cm-siRNAs.



**Figure S5.** Cell lines distribution in siRNAmoD database.