

Supplemental Fig -1

CELLO RESULTS

SeqID: unknow

Analysis Report:

SVM	LOCALIZATION	RELIABILITY
Amino Acid Comp.	Periplasmic	0.262
N-peptide Comp.	Cytoplasmic	0.715
Partitioned seq. Comp.	Cytoplasmic	0.387
Physico-chemical Comp.	Cytoplasmic	0.630
Neighboring seq. Comp.	Cytoplasmic	0.328

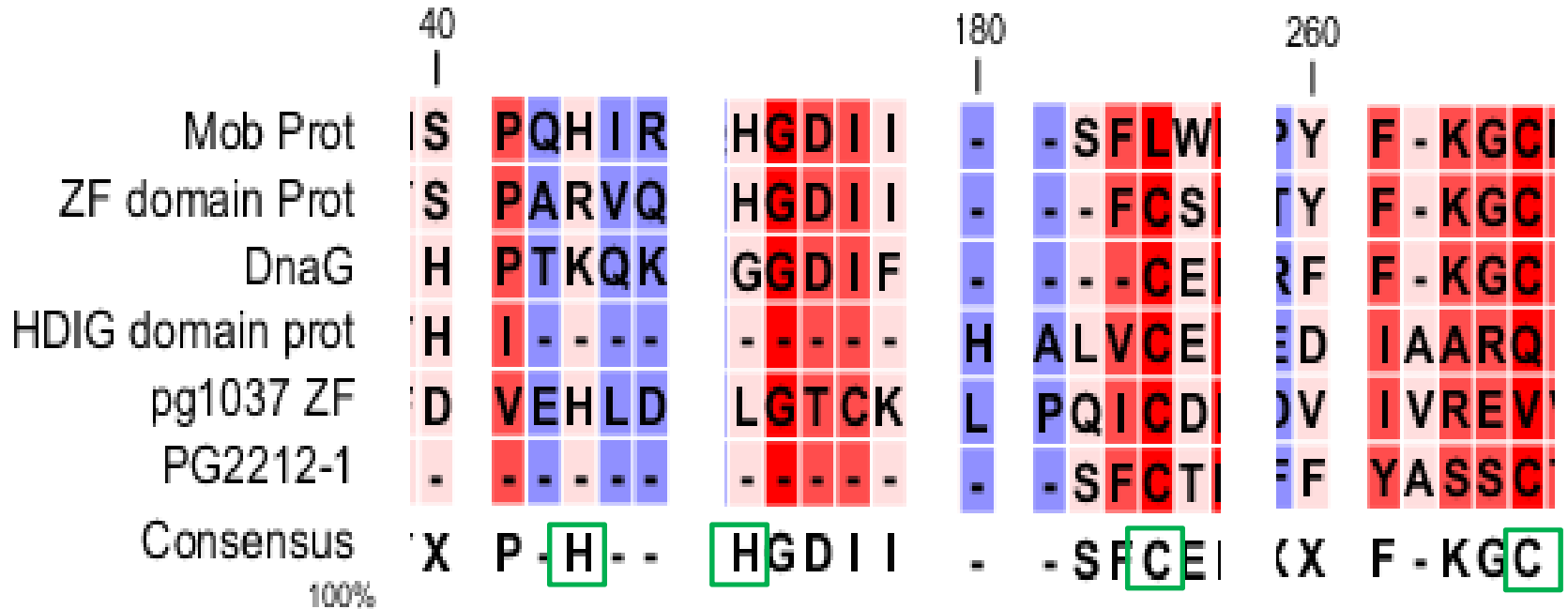
CELLO Prediction:

Cytoplasmic	2.299 *
Periplasmic	1.022
Extracellular	0.726
OuterMembrane	0.502
InnerMembrane	0.451

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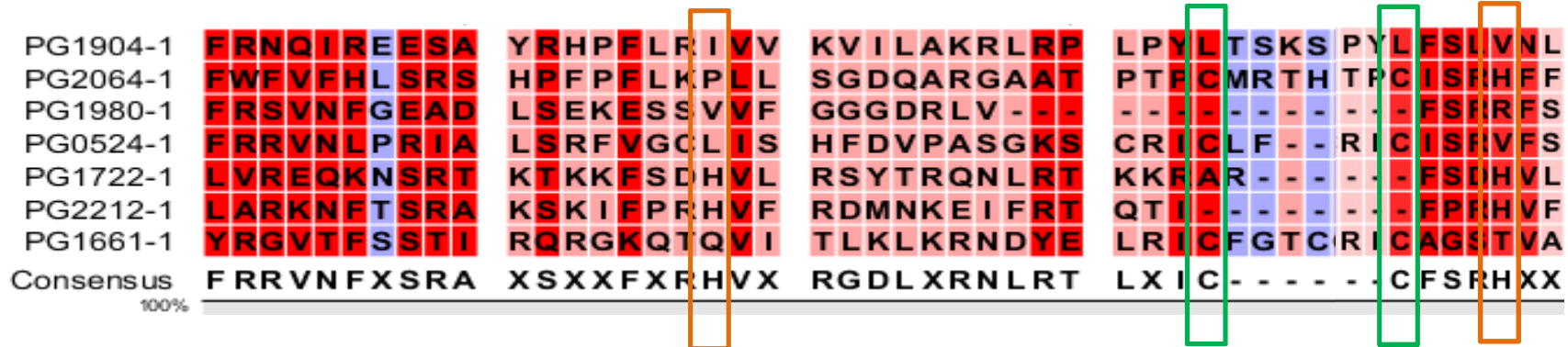
In silico sub-cellular localization of PG2212 protein was predicted to be cytoplasmic.

Supplemental Fig -2



Multiple sequence alignment of *P. gingivalis* proteins containing zinc finger domains.

Supplemental Fig -3



Multiple sequence alignment of 7 *P. gingivalis* proteins (PG2212, PG2064, PG1980, PG1904, PG1722, PG1661 And PG0524) showing conserved DUF 1661 domain and zinc finger target sites.