

Gene Cluster ID	Atp6v1H
Drosophila melanogaster gene	VhaSFD (CG17332)
FlyBase ID	FBgn0027779
Predicted function	V-type ATPase, H⁺ transporting, V1 subunit H
Cytogenetic map	2L:6A12-36A12
Gene sequence location	2L:16723138..16727825 [-]
Dmel CLEAR sequence	TCACAAGAccagTCAGCTGA
CLEAR sequence location	2L:16727801..16727782
FlyBase VhaSFD (CG17332) gene model & CLEAR element (red) positions:	



Atp6v1H (VhaSFD) gene structure comparison:

Key: UTR region, CDS region, intron, alternatively spliced exon

