

Scale chr4: 307,500 | 308,000 | 308,500 | 309,000 | 309,500 | 310,000 | 310,500 | 311,000 | 311,500

FBS predictions in Drosophila melanogaster (dm6) for all profiles in the JASPAR CORE insects collection (2020) (11981 items filtered out)

Gene	307,500	308,000	308,500	309,000	309,500	310,000	310,500	311,000	311,500
HEX	Ubx	inv	slp1	hb	Clamp	Awh	C15	CG11294	CG32105
CG11294	CG32105	CG32532	CG11294	CG32105	CG32532	CG11617	br(var.4)	vfi	
HGTX	Lim1	Lim3	HGTX	Lim1	Lim3	PHDP	Vsx1	abd-A	
PHDP	abd-A								
HEX	CG11617								

FBS predictions in Drosophila melanogaster (dm6) for all profiles in the JASPAR CORE insects collection (2022) (12108 items filtered out)

Gene	307,500	308,000	308,500	309,000	309,500	310,000	310,500	311,000	311,500
HEX	Ubx	inv	slp1	hb	Clamp	Awh	C15	CG11294	CG32105
CG11294	CG32105	CG32532	CG11294	CG32105	CG32532	CG11617	dsx	pan	br
HGTX	Lim1	Lim3	HGTX	Lim1	Lim3	PHDP	Vsx1	abd-A	
PHDP	abd-A								
HEX	CG11617								

Gap Locations

