

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	Ubx	Antp	Ubx	Stat92E
CG11294	C15	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E
CG32105	C15	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E
CG32532	C15	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E
HGTX	Lim1	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3
PHDP	Lim1	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3
abd-A	al	en	hbn	lab	otp	pb	repo	unpg	zen2
HEX	Ubx	inv	slp1	hb	Clamp	Ubx	Antp	Ubx	Stat92E
CG11617	C15	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E

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	Ubx	Antp	Ubx	Stat92E
CG11294	C15	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E
CG32105	C15	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E
CG32532	C15	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E
HGTX	Lim1	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3
PHDP	Lim1	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3	Lim3
abd-A	al	en	hbn	lab	otp	pb	repo	unpg	zen2
HEX	Ubx	inv	slp1	hb	Clamp	Ubx	Antp	Ubx	Stat92E
CG11617	C15	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E	Stat92E

