

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	Awh	C15	CG11294	CG32105	CG32532	CG11294
CG32105	CG32532	CG11294	CG32105	CG32532	CG11294	CG32105	CG32532	CG11294	CG32105
CG32532	E5	H2.0	HGTX	Lim1	Lim3	PHDP	Vsx1	abd-A	al
	en	hbn	lab	otp	repo	unpg	zen2	Cf2	Dbx
	HEX	CG11617	br(var.4)	Stat92E	onecut				

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	Awh	C15	CG11294	CG32105	CG32532	CG11294
CG32105	CG32532	CG11294	CG32105	CG32532	CG11294	CG32105	CG32532	CG11294	CG32105
CG32532	Vsx1	E5	HGTX	Lim1	Lim3	PHDP	Vsx1	abd-A	al
	en	hbn	lab	otp	repo	unpg	zen2	H2.0	PHDP
	HEX	CG11617	br(var.4)	Stat92E	onecut				

