

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	Scr	Dbx
CG11294	C15	Stat92E	Stat92E	Awh	C15	Antp	Dfd	ems	Ubx
CG32105		Stat92E	Stat92E	C15	ems	Ubx	inv	br	
CG32532			CG11294	CG32105	CG32532	CG11617	br(var.4)	inv	
HGTX	Lim1	Lim3	HGTX	Lim1	Lim3	PHDP	Vsx1	abd-A	
PHDP	abd-A		PHDP	Lim1	Lim3	PHDP	Vsx1	abd-A	
HEX	CG11617		CG11294	CG32105	CG32532	CG11294	CG32105	CG32532	
CG11617			CG11294	CG32105	CG32532	CG11294	CG32105	CG32532	

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	Scr	Dbx
CG11294	C15	Stat92E	Stat92E	Awh	C15	Antp	Dfd	ems	Ubx
CG32105		Stat92E	Stat92E	C15	ems	Ubx	inv	br	
CG32532			CG11294	CG32105	CG32532	CG11617	br(var.4)	inv	
HGTX	Lim1	Lim3	HGTX	Lim1	Lim3	PHDP	Vsx1	abd-A	
PHDP	abd-A		PHDP	Lim1	Lim3	PHDP	Vsx1	abd-A	
HEX	CG11617		CG11294	CG32105	CG32532	CG11294	CG32105	CG32532	
CG11617			CG11294	CG32105	CG32532	CG11294	CG32105	CG32532	

