

Supplementary Table 6: *In silico* canSNP typing in intragenic region of *Coxiella burnetii* isolates sequenced in this study.

Isolate name	Host	Country	CanSNP in intragenic conserved region										Derived for CanSNP ^b	SNP type
			C.1	C.3	C.4	C.5	C.6	C.7	C.8	C.9	C.10	C.11		
Nine Mile	Tick	US	C	T	G	T	G	A	C	A	T	G	C10	3
CbBEC1	Cattle	Belgium	C	NA ^c	G	T	G	A	C	A	NA ^c	A	C11	1
CbBEB1	Cattle	Belgium	C	T	G	T	G	A	C	A	C	A	C11	2
CbBEB2	Goat	Belgium	C	T	G	T	G	A	C	A	C	NA ^c	C11	2
CbBEC2	Goat	Belgium	C	T	G	T	G	A	C	A	C	A	C11	2
Henzerling ^a	Human	Italy	C	T	G	T	G	A	C	A	C	A	C11	8

^a Genetic profile from *Karlsson et al.* (2014) for comparison with isolates from this study.

^b Based on phylogeny from *Karlsson et al.* (2014).

^c NA: No available result from MLVA typing.