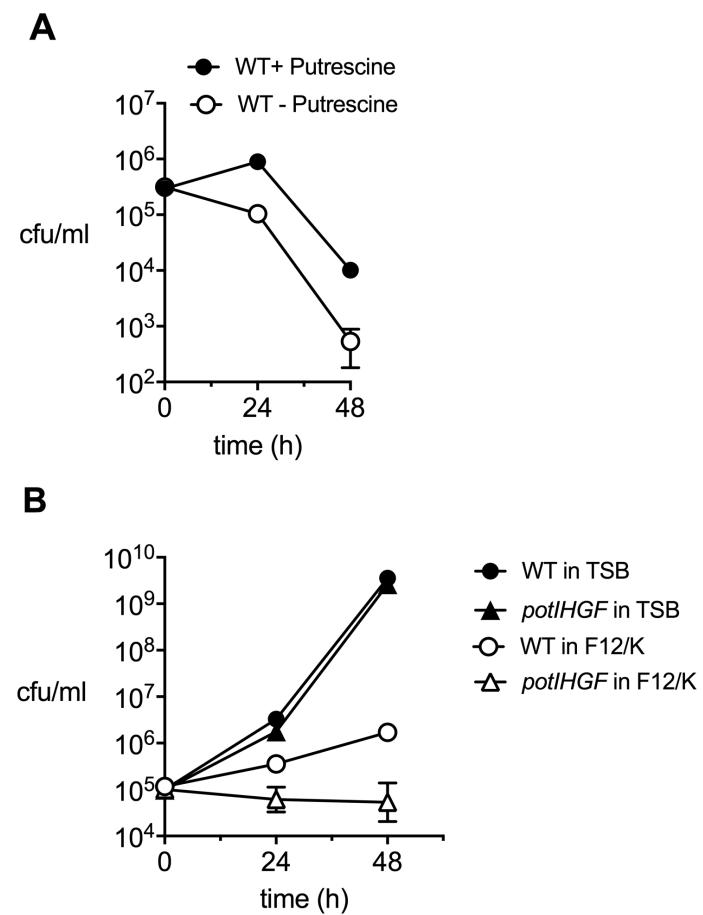


Supplementary information for Kerrinnes et al., **Utilization of host polyamines in alternatively activated macrophages promotes chronic infection by *Brucella abortus***

Fig. S1

Table S1. Composition of modified F12/K media

Table S2. Real-time PCR oligonucleotides



**Fig. S1:** (A) Growth of *B. abortus* in F12/K medium formulated with or without putrescine. (B) Growth of *B. abortus* 2308 or *potlHGF* mutant in TSB or in F12/K medium containing putrescine as sole carbon source.

**Table S1. Composition of modified F12/K media**

	concentration [mg/L]	source
<b>amino acids</b>		
MEM Amino Acids Solution	1X	Life Technologies
MEM Non-Essential Amino Acids Solution	1X	Life Technologies
L-Glutamine	292	Life Technologies
<b>Vitamins</b>		
1X MEM Vitamin Solution		Life Technologies
Biotin	0.07	Sigma-Aldrich
Vitamin B <sub>12</sub>	1.4	Sigma-Aldrich
<b>Inorganic Salts and additional substances</b>		
Calcium Chloride (CaCl <sub>2</sub> ) (anhyd.)	102	Sigma-Aldrich
Cupric sulfate (CuSO <sub>4</sub> -5H <sub>2</sub> O)	0.002	Sigma-Aldrich
Ferric sulfate (FeSO <sub>4</sub> -7H <sub>2</sub> O)	0.8	Sigma-Aldrich
Magnesium Chloride (anhyd.)	49.7	Sigma-Aldrich
Magnesium Sulfate (MgSO <sub>4</sub> ) (anhyd.)	192	Sigma-Aldrich
Sodium Chloride (NaCl)	8500	Sigma-Aldrich
Sodium Bicarbonate (NaHCO <sub>3</sub> )	2500	Sigma-Aldrich
Sodium Chloride (NaCl)	7530	Sigma-Aldrich
Sodium Phosphate dibasic (Na <sub>2</sub> HPO <sub>4</sub> ) (anhyd.)	115.5	Sigma-Aldrich
Sodium Phosphate monobasic (NaH <sub>2</sub> PO <sub>4</sub> ) (anhyd.)	59	Sigma-Aldrich
Zinc sulfate (ZnSO <sub>4</sub> -7H <sub>2</sub> O)	0.144	Sigma-Aldrich
D-Glucose (Dextrose)	1260	Sigma-Aldrich
Hypoxanthine Na	4	Sigma-Aldrich
Lipoic Acid	0.21	Sigma-Aldrich
Phenol Red	3	Sigma-Aldrich
Sodium Pyruvate	220	Sigma-Aldrich
Thymidine	0.7	Sigma-Aldrich
Putrescine 2HCl, when needed	0.32	Santa Cruz - Biotechnology

**Table S2. Real-time PCR oligonucleotides used in the study:**

<b>primer sequence (5' to 3')</b>	
mArg1-Fwd	GCTGTCTTCCCAAGAGTTGGG
mArg1-Rev	ATGGAAGAGACCTTCAGCTAC
mRplp0-2-Fwd	GATGGGCAACTGTACCTGACTG
mRplp0-2-Rev	CTGGGCTCCTCTTGGAAATG
mSlc7a2-Fwd	GCCTTGTTGGGCTTGACTG
mSlc7a2-Rev	CCATCCTCCGCCATAGCATA
mYm1-Fwd	GGGCATACCTTATCCTGAG
mYm1-Rev	CCACTGAAGTCATCCATGTC