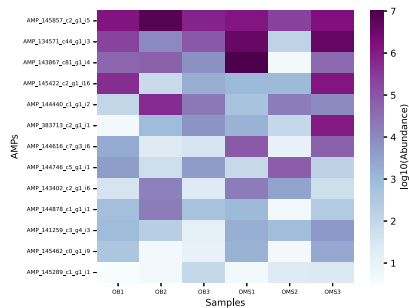


# AMPs present in all samples

## Heatmap



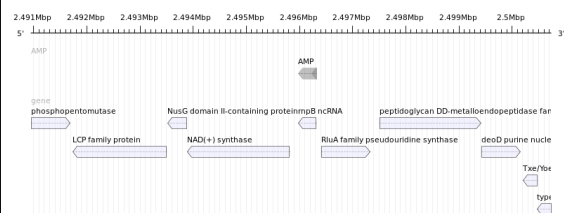
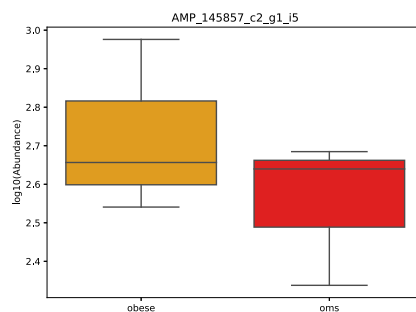
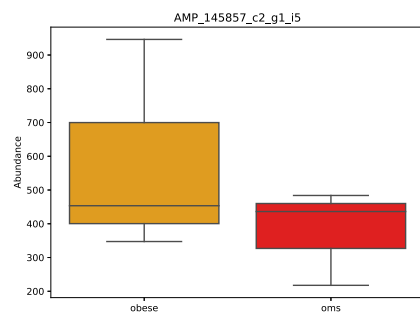
## Classification and secueneces

Transcript	Best-hit	AMP	Sequence
TRINITY_DN145857_c2_g1_i5	Faecalibacterium prausnitzii	AMP_145857_c2_g1_i5.p4	ENEWNIRRRGGFDRSEESQGLTEHGNC
TRINITY_DN134571_c44_g1_i3	Not assigned	AMP_134571_c44_g1_i3.p7	MGSTPIASTNM
TRINITY_DN143867_c81_g1_i4	Eubacteriales	AMP_143867_c81_g1_i4.p7	GQDAG
TRINITY_DN145422_c2_g1_i16	Faecalibacterium prausnitzii	AMP_145422_c2_g1_i16.p11	MKRVPQPEK
TRINITY_DN144440_c1_g1_i2	Not assigned	AMP_144440_c1_g1_i2.p8	MGSTPIAST
TRINITY_DN383713_c2_g1_i1	Not assigned	AMP_383713_c2_g1_i1.p8	MGSTPIASTND
TRINITY_DN144616_c7_g3_i6	Eubacteriales	AMP_144616_c7_g3_i6.p5	GDANPRLVSCGAALYKWR
TRINITY_DN144746_c5_g1_i1	Clostridium	AMP_144746_c5_g1_i1.p8	MRGHTGLMSSVKLSC
TRINITY_DN143402_c2_g1_i6	Eubacteriales	AMP_143402_c2_g1_i6.p8	EESPGFAGQAAG
TRINITY_DN144878_c1_g1_i1	Faecalibacterium prausnitzii	AMP_144878_c1_g1_i1.p9	KKKGAKTEG
TRINITY_DN141259_c3_g4_i3	Eubacteriales	AMP_141259_c3_g4_i3.p6	MQQH
TRINITY_DN145462_c0_g1_i9	Faecalibacterium prausnitzii	AMP_145462_c0_g1_i9.p9	MGDYAAL
TRINITY_DN145289_c1_g1_i1	Faecalibacterium prausnitzii	AMP_145289_c1_g1_i1.p7	MGGYAAL

## Individual boxplots

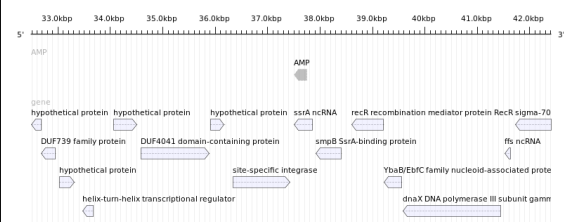
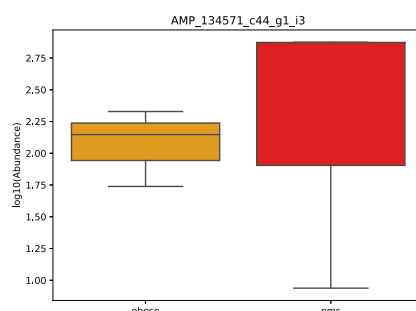
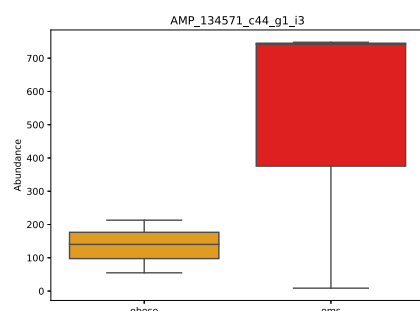
### AMP\_145857\_c2\_g1\_i5.p4

Transcript	Best-hit	AMP	Sequence
TRINITY_DN145857_c2_g1_i5	Faecalibacterium prausnitzii	AMP_145857_c2_g1_i5.p4	ENEWNIRRRGGFDRSEESQGLTEHGNC



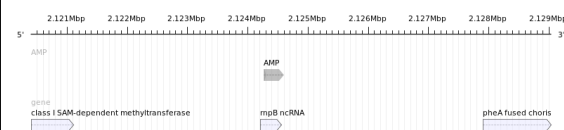
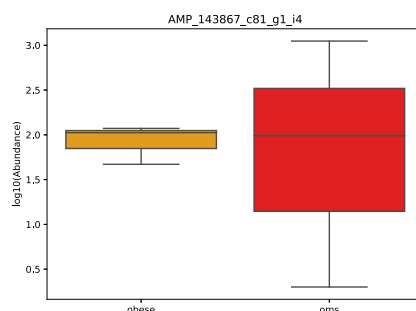
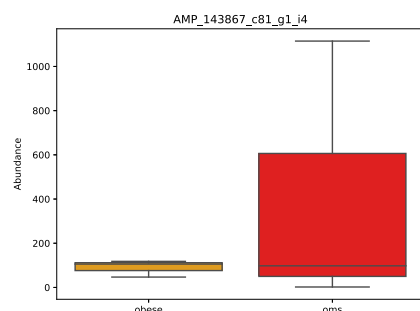
## AMP\_134571\_c44\_g1\_i3.p7

Transcript	Best-hit	AMP	Sequence
TRINITY_DN134571_c44_g1_i3	Not assigned	AMP_134571_c44_g1_i3.p7	MGSTPIASTNM



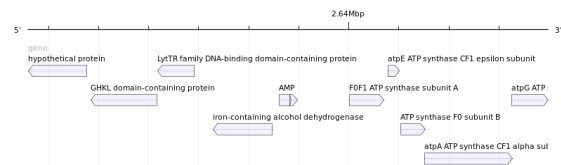
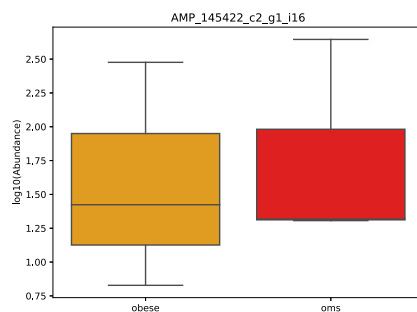
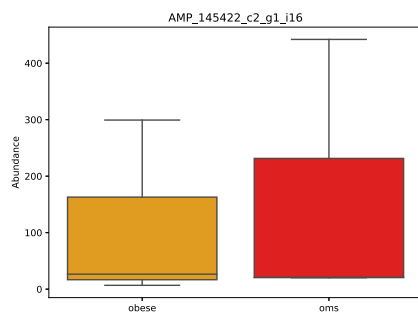
## AMP\_143867\_c81\_g1\_i4.p7

Transcript	Best-hit	AMP	Sequence
TRINITY_DN143867_c81_g1_i4	Eubacteriales	AMP_143867_c81_g1_i4.p7	GQDAG



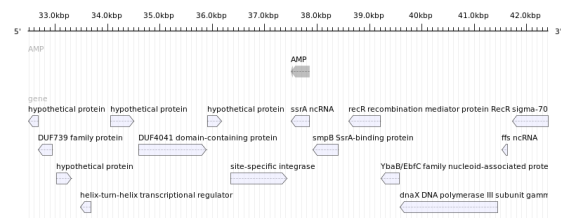
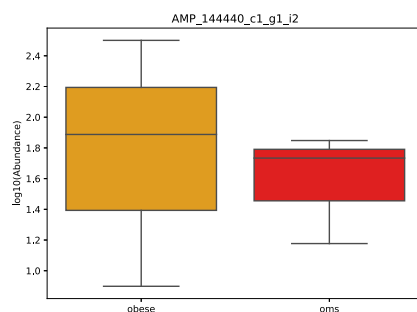
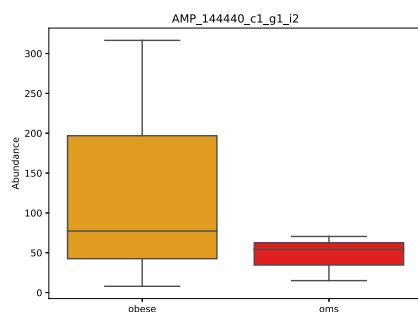
## AMP\_145422\_c2\_g1\_i16.p11

Transcript	Best-hit	AMP	Sequence
TRINITY_DN145422_c2_g1_i16	Faecalibacterium prausnitzii	AMP_145422_c2_g1_i16.p11	MKRPVQPEK



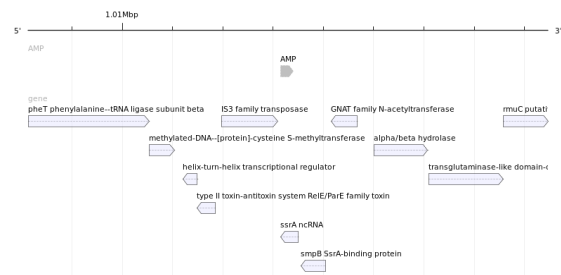
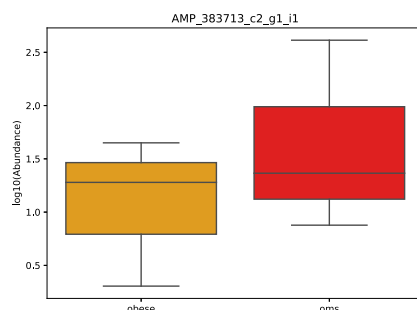
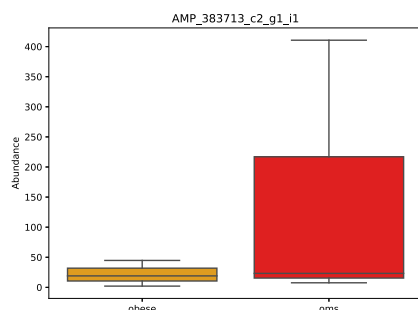
## AMP\_144440\_c1\_g1\_i2.p8

Transcript	Best-hit	AMP	Secuence
TRINITY_DN144440_c1_g1_i2	Not assigned	AMP_144440_c1_g1_i2.p8	MGSTPIAST



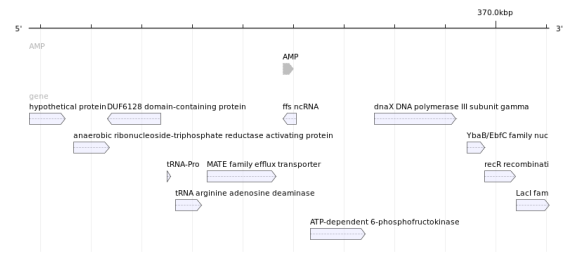
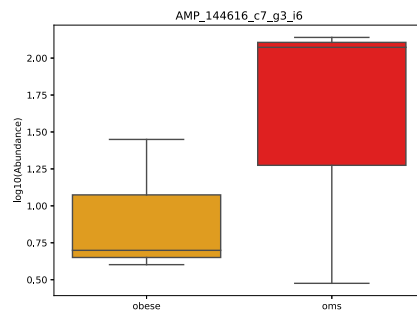
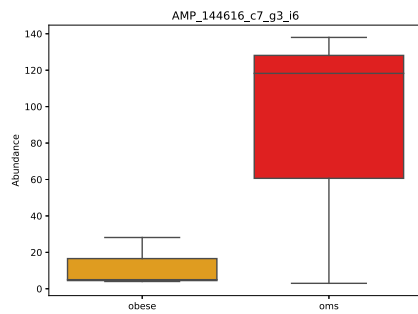
## AMP\_383713\_c2\_g1\_i1.p8

Transcript	Best-hit	AMP	Secuence
TRINITY_DN383713_c2_g1_i1	Not assigned	AMP_383713_c2_g1_i1.p8	MGSTPIASTND



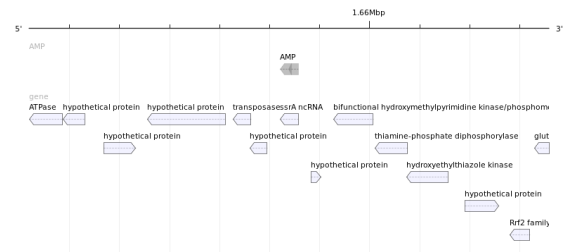
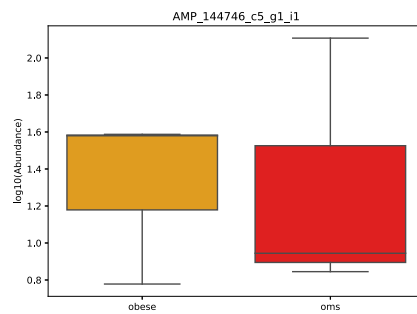
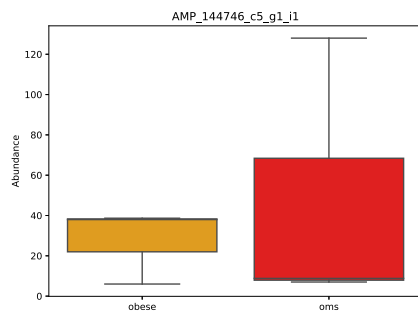
## AMP\_144616\_c7\_g3\_i6.p5

Transcript	Best-hit	AMP	Secuence
TRINITY_DN144616_c7_g3_i6	Eubacteriales	AMP_144616_c7_g3_i6.p5	GDANPRLVSCGAALYKWR



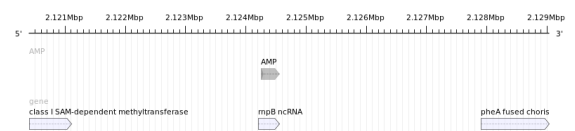
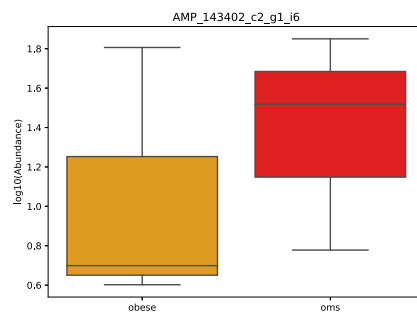
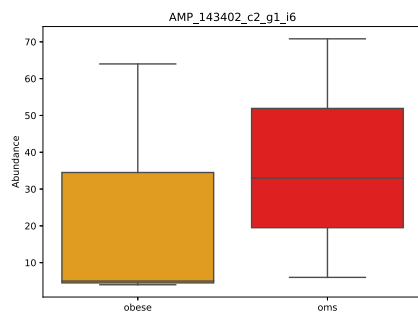
## AMP\_144746\_c5\_g1\_i1.p8

Transcript	Best-hit	AMP	Sequence
TRINITY_DN144746_c5_g1_i1	Clostridium	AMP_144746_c5_g1_i1.p8	MRGHTGLMSSVKLSC



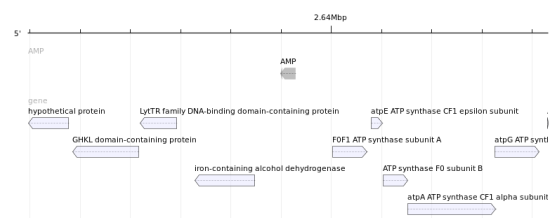
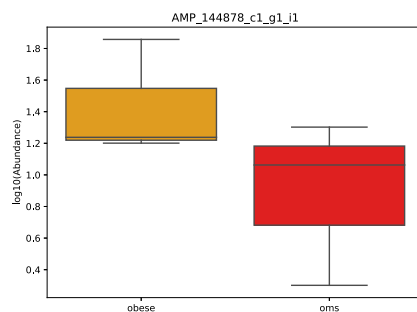
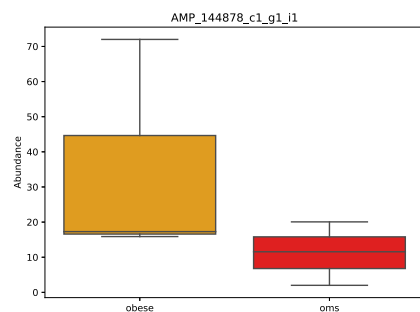
## AMP\_143402\_c2\_g1\_i6.p8

Transcript	Best-hit	AMP	Sequence
TRINITY_DN143402_c2_g1_i6	Eubacteriales	AMP_143402_c2_g1_i6.p8	EESPGFAGQAAG



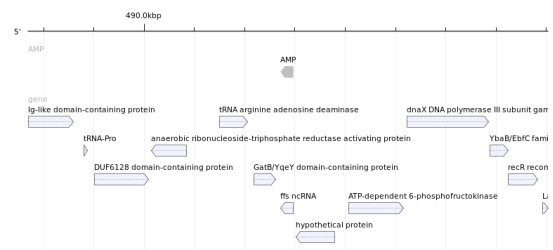
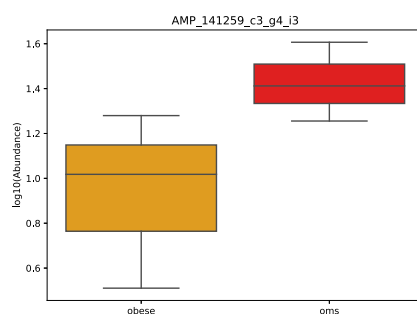
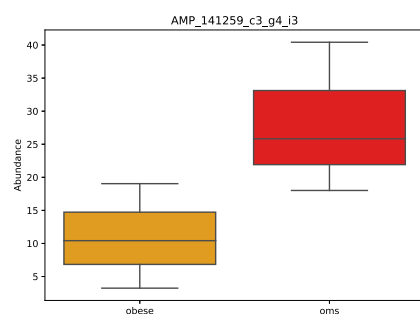
## AMP\_144878\_c1\_g1\_i1.p9

Transcript	Best-hit	AMP	Sequence
TRINITY_DN144878_c1_g1_i1	Faecalibacterium prausnitzii	AMP_144878_c1_g1_i1.p9	KKKGAKTEG



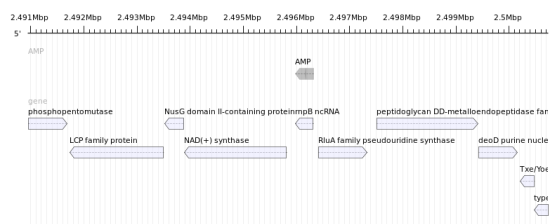
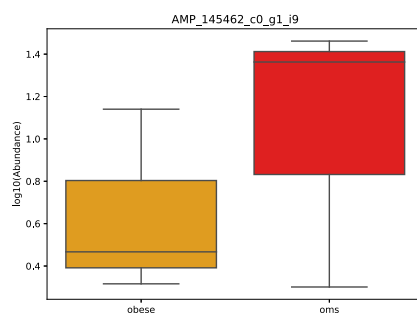
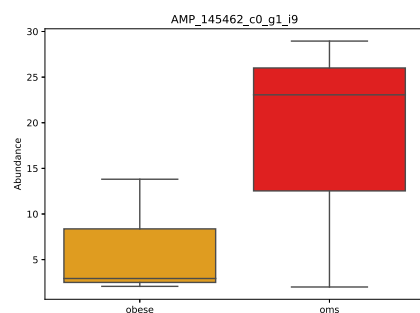
## AMP\_141259\_c3\_g4\_i3.p6

Transcript	Best-hit	AMP	Sequence
TRINITY_DN141259_c3_g4_i3	Eubacteriales	AMP_141259_c3_g4_i3.p6	MQQH



## AMP\_145462\_c0\_g1\_i9.p9

Transcript	Best-hit	AMP	Sequence
TRINITY_DN145462_c0_g1_i9	Faecalibacterium prausnitzii	AMP_145462_c0_g1_i9.p9	MGDYAAL



## AMP\_145289\_c1\_g1\_i1.p7

Transcript	Best-hit	AMP	Sequence
TRINITY_DN145289_c1_g1_i1	Faecalibacterium prausnitzii	AMP_145289_c1_g1_i1.p7	MGGYAAL

