

mRNA	Protein	
1.Ribosome <input type="checkbox"/> 2.Purine metabolism <input type="checkbox"/>		lowMg
1.Flagellar assembly <input type="checkbox"/> 2.Sulfur metabolism <input type="checkbox"/> 3.Nitrogen metabolism <input type="checkbox"/>	1.Biosynthesis of siderophore group nonribosomal peptides <input type="checkbox"/> 2.Two-component system <input type="checkbox"/> 3.Pyruvate metabolism <input type="checkbox"/>	highMg
1.Flagellar assembly <input type="checkbox"/>	1.Biosynthesis of amino acids <input type="checkbox"/> 2.Biosynthesis of secondary metabolites <input type="checkbox"/> 3.Biosynthesis of antibiotics <input type="checkbox"/> 4.Metabolic pathways <input type="checkbox"/> 5.Phenylalanine, tyrosine and tryptophan biosynthesis <input type="checkbox"/>	highNa
1.Ribosome <input type="checkbox"/> 2.Biosynthesis of antibiotics <input type="checkbox"/>		glycerol
1.Pentose phosphate pathway <input type="checkbox"/>	1.Biosynthesis of siderophore group nonribosomal peptides <input type="checkbox"/>	gluconate
1.Pyruvate metabolism <input type="checkbox"/> 2.Ribosome <input type="checkbox"/>	1.Citrate cycle (TCA cycle) <input type="checkbox"/> 2.Pyruvate metabolism <input type="checkbox"/> 3.Carbon metabolism <input type="checkbox"/>	lactate