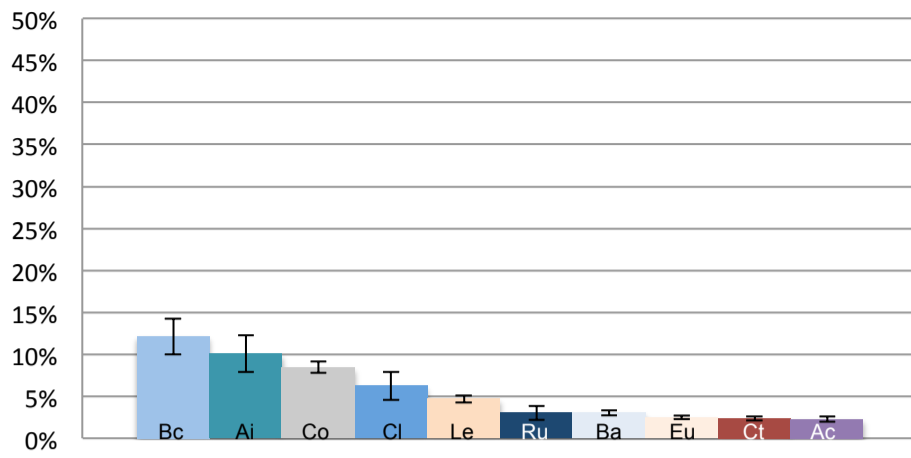
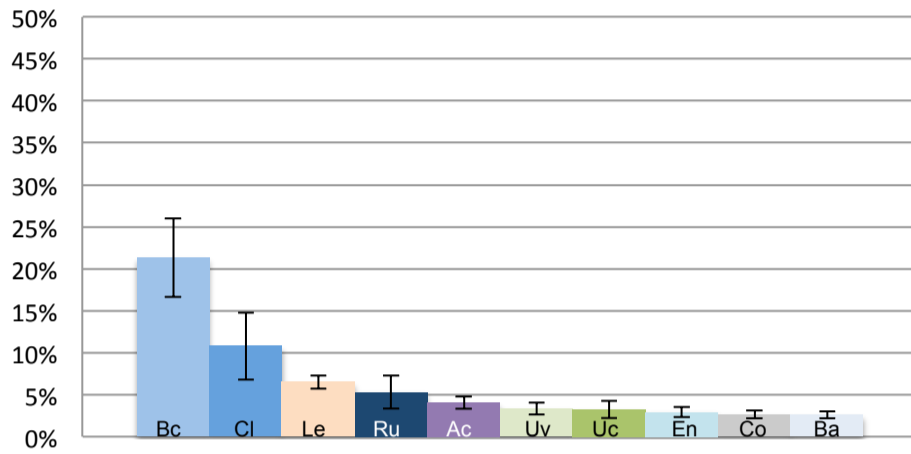


All samples (n=12)

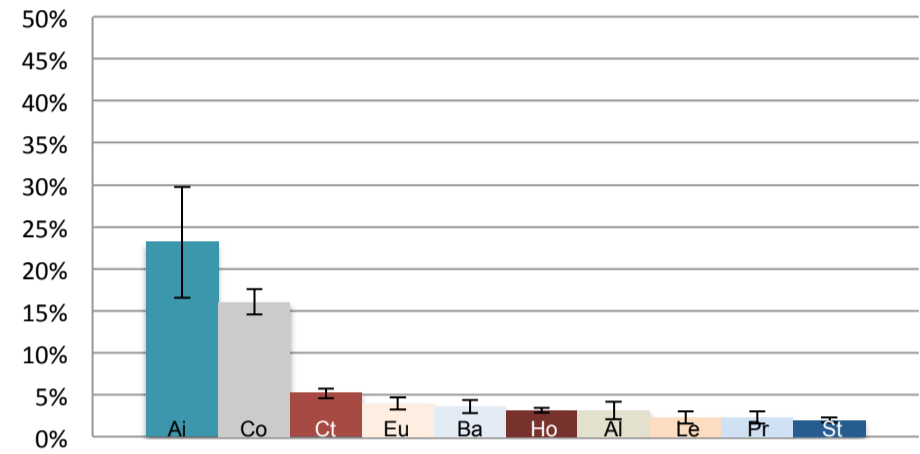


cellular organisms	(Co)	Bacteria	(Ba)
Eukaryota	(Eu)	Proteobacteria (Bac)	(Pr)
Acanthamoeba (Euk, Amoebozoa)	(Ac)	Gamma proteobacteria (Bac)	(Ga)
Leotiomyceta (Euk, Fungi)	(Le)	Enterobacteriaceae (Bac)	(En)
Exophiala (Euk, Fungi)	(Ex)	Diploricetisia (Bac)	(Di)
Mycosphaerella (Euk, Fungi)	(My)	Rickettsiella (Bac)	(Ri)
Claviceps (Euk, Fungi)	(Ca)	Acinetobacter (Bac)	(Ai)
Boreoeutheria (Euk, Fungi)	(Bo)	Pseudomonas (Bac)	(Ps)
Simiiformes (Euk, Primates)	(Si)	Alphaproteobacteria (Bac)	(Al)
Catarrhini (Euk, Primates)	(Ct)	Bacillus (Bac)	(Bc)
Homo (Euk, Primates)	(Ho)	Clostridiales (Bac)	(Cl)
unclassified viruses	(Uv)	Ruminiclostridium (Bac)	(Ru)
unclassified Circoviridae (Vir)	(Uc)	Staphylococcus (Bac)	(St)
		Gardnerella (Bac)	(Gr)
		Propionibacterium (Bac)	(Pp)

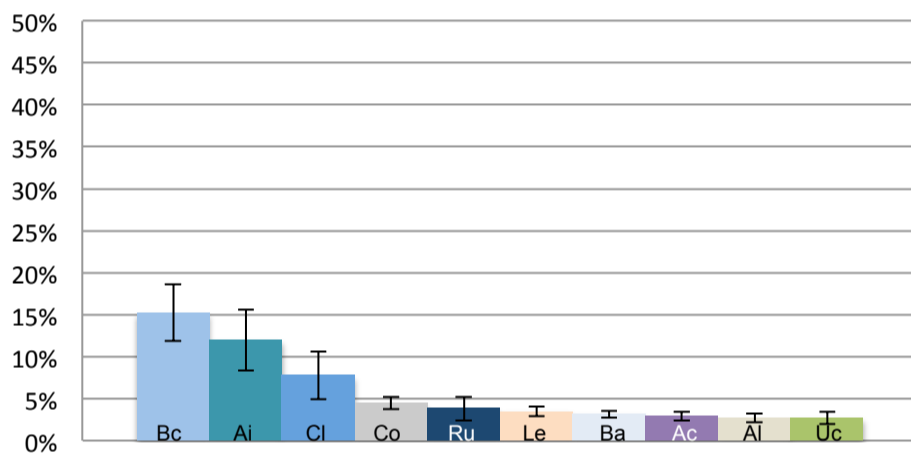
PMA: SAF + GA (n=6)



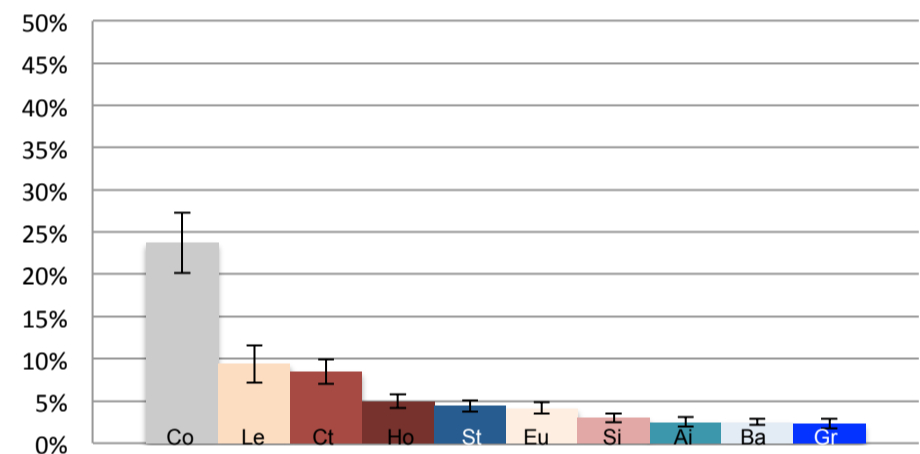
nonPMA: SAF + GA (n=6)



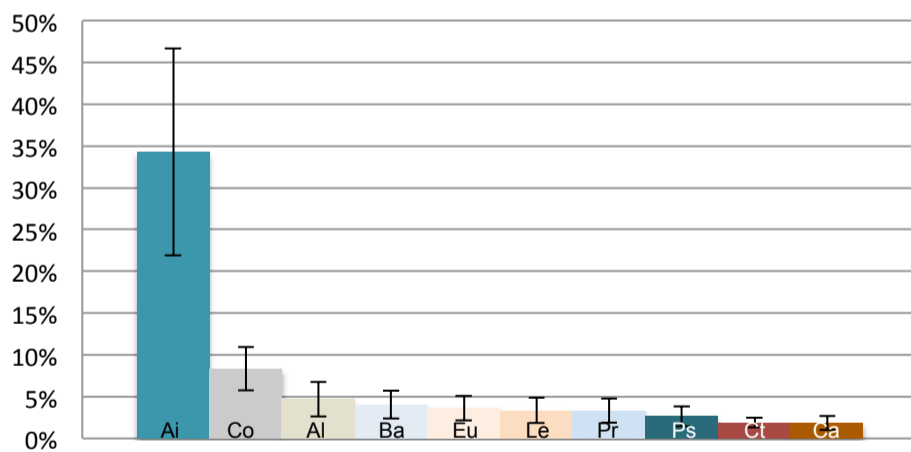
SAF: PMA + nonPMA (n=6)



GA: PMA + nonPMA (n=6)



SAF_nonPMA (n=3)



GA_nonPMA (n=3)

