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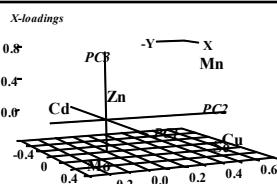
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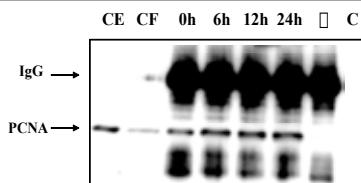
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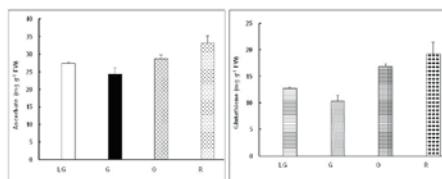
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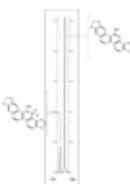
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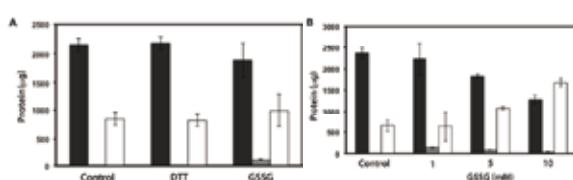
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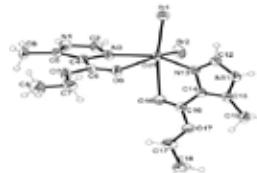
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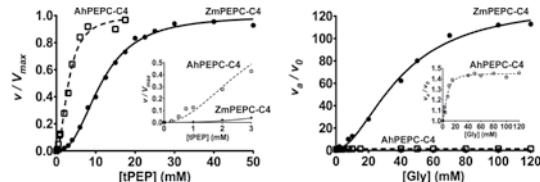
Table 1. Proteins identified by mass spectrometry as components of Cap-binding complexes in maize non-germinated (0h) or germinated (24h) seeds.

Band ID	Protein ID	NCBI Acc #	MW	MW _{obs}	Number of peptides ^a	% Protein Coverage
FT1	Heat shock protein HSP101	GI4584957	101	100	10	-
	Component of the nucleic acid binding complex (CBC), CBP80	GI108708115	59	100	4	4
	Component of the eukaryotic translation initiation factor 3 complex, eIF3c	GI113535264	104	100	3	-

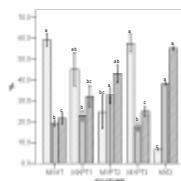
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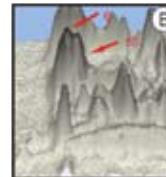
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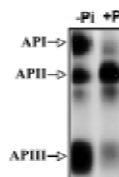
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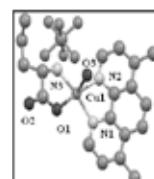
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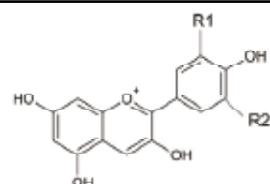


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César Aza-González, Héctor Gordon Núñez-Palenius, and Neftalí Ochoa-Alejo*



Anthocyanin	R1	R2
Cyanidin	OH	H
Delphinidin	OH	OH
Malvidin	O-CH ₃	O-CH ₃
Petunidin	H	H
Peonidin	O-CH ₃	H
Pelargonidin	O-CH ₃	OH