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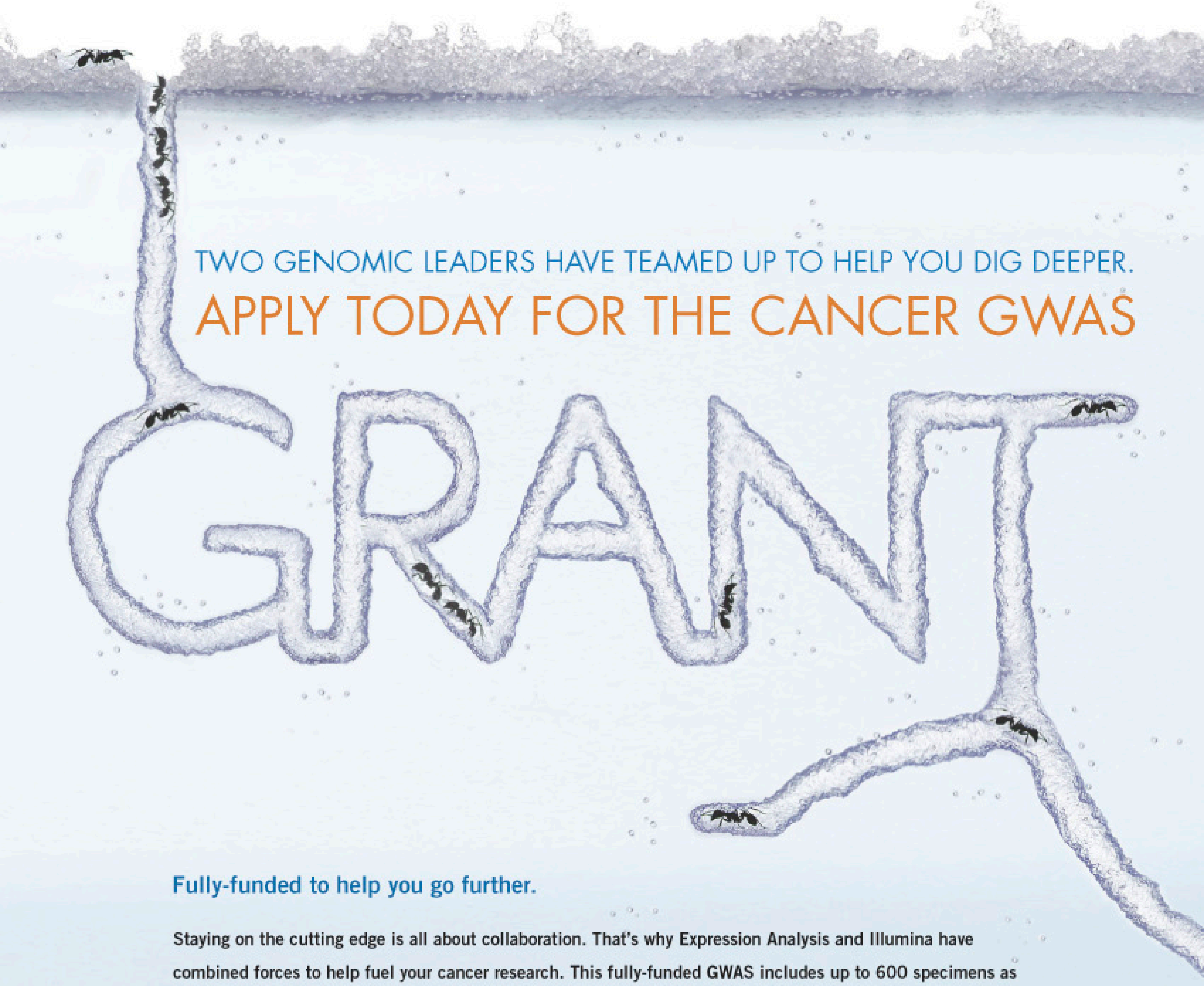

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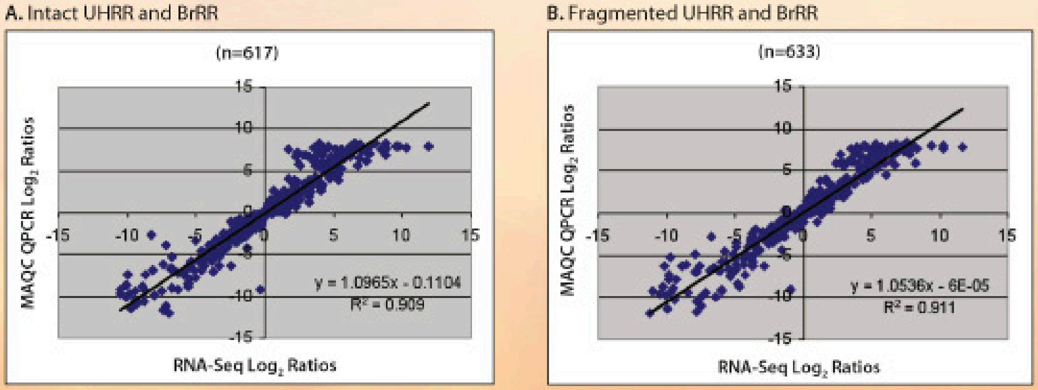
A. Intact UHRR and BrRR
(n=617)
15
10
5
-15
-10
-5
0
5
10
15
-5
-10
-15
MAQC QPCR Log₂ Ratios
RNA-Seq Log₂ Ratios
y = 1.0965x - 0.1104
R² = 0.909
B. Fragmented UHRR and BrRR
(n=633)
15
10
5
-15
-10
-5
0
5
10
15
-5
-10
-15
MAQC QPCR Log₂ Ratios
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y = 1.0536x - 6E-05
R² = 0.911

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- Cassava Genomics
- Cattle/Sheep/Swine
- Citrus Genome
- Coffee
- Comparative Genomics
- Cool Season Legumes
- Cotton Genome Initiative
- Cucurbit
- Ecological Genomics
- Equine
- Evolution of Genome Size
- Forage & Turf Plants
- Forest Tree
- Fruit/Nuts
- Functional Genomics
- Fungal Genomics
- Gene Expression Analysis
- Genomics Assisted Breeding
- Genomics for Disease Resistance
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- Grass Genome Initiative
- Host-Microbe Interactions
- Large Insert DNA Library
- Lolium Genome Initiative
- Maize
- Molecular Markers
- Mutation Screening
- Oats
- Ornamentals
- Phylogenomics
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- Poultry
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