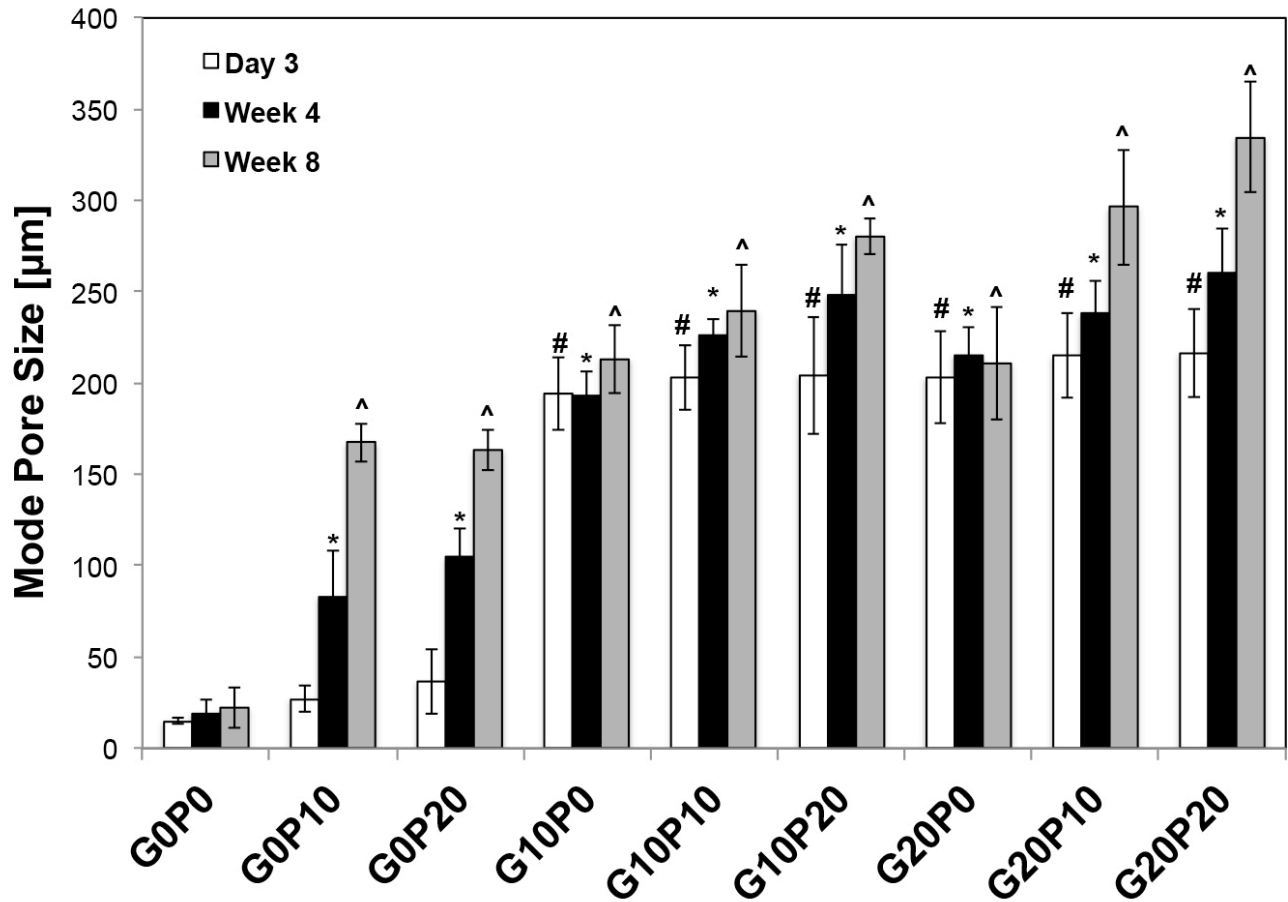


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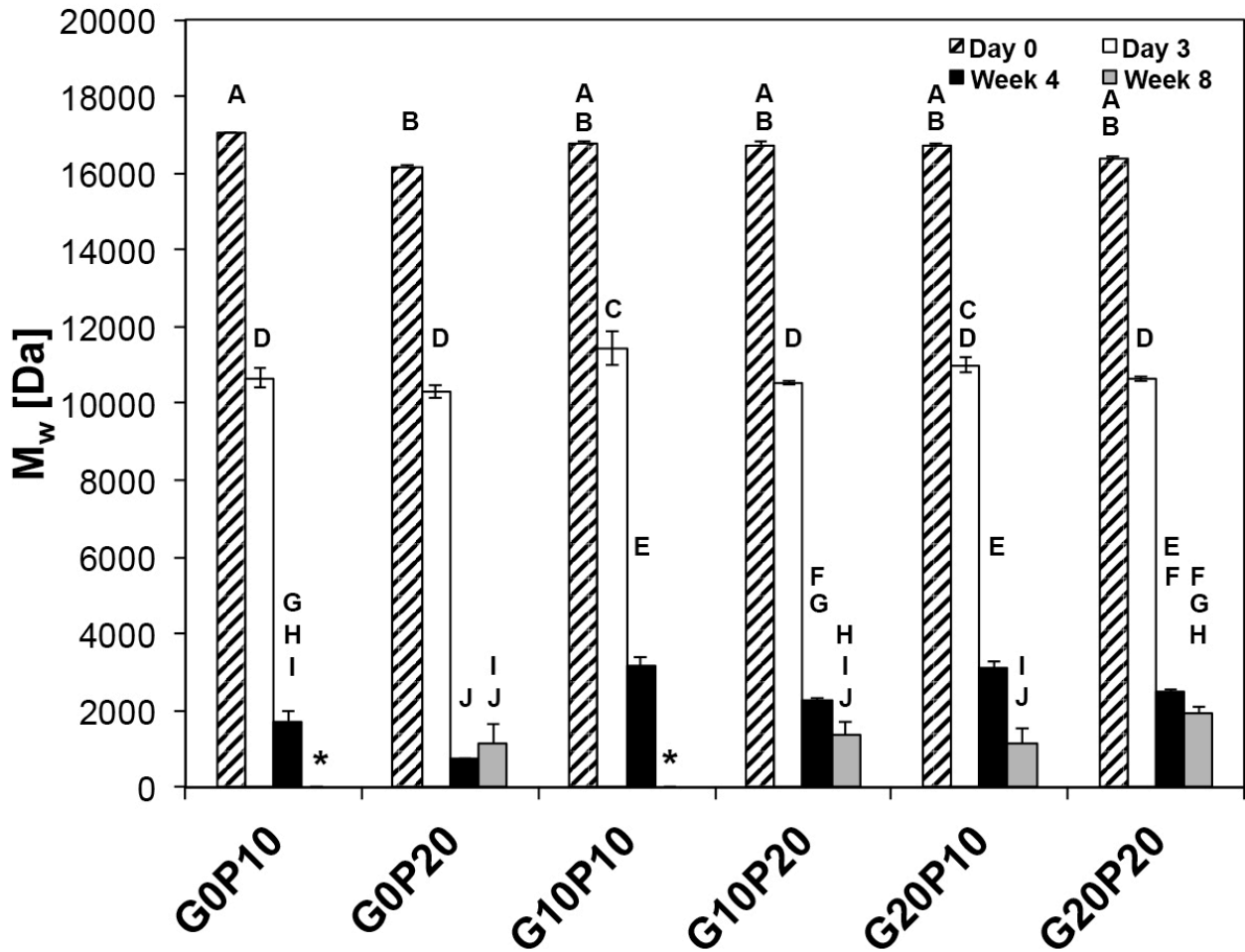
## Supplemental Figures and Tables

Figure S1



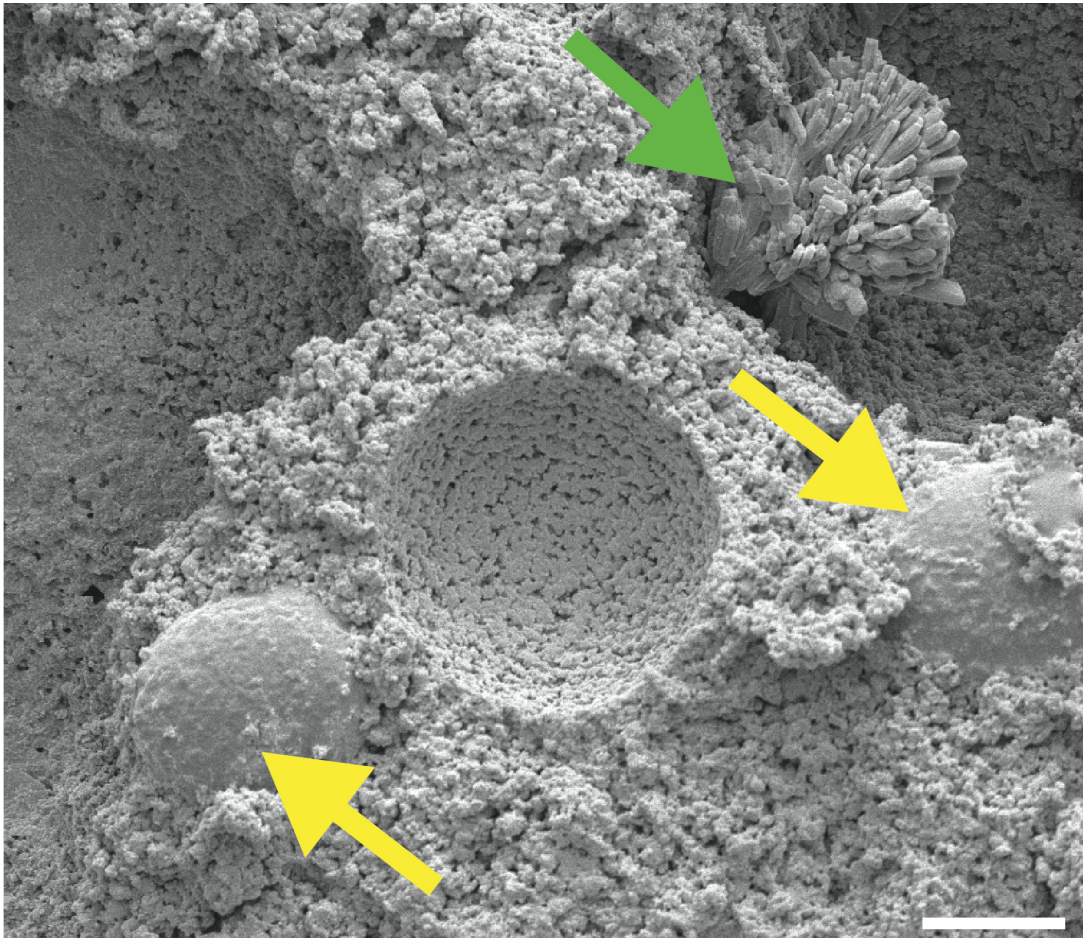
**Figure S1.** Mode pore size of the GMP/PLGA/CPC composites determined by microCT after being leached in PBS at 37°C for 3 days, 4 wks, and 8 wks. GMPs and PLGA MPs were introduced in three different weight fractions (0, 10, and 20%). Within the Day 3 series, # denotes significance with respect to the CPC control. In the Week 4 series, \* denotes significance with respect to the CPC control. In the Weeks 8 series, ^ denotes significance with respect to the control (n=3, p<0.05).

**Figure S2**



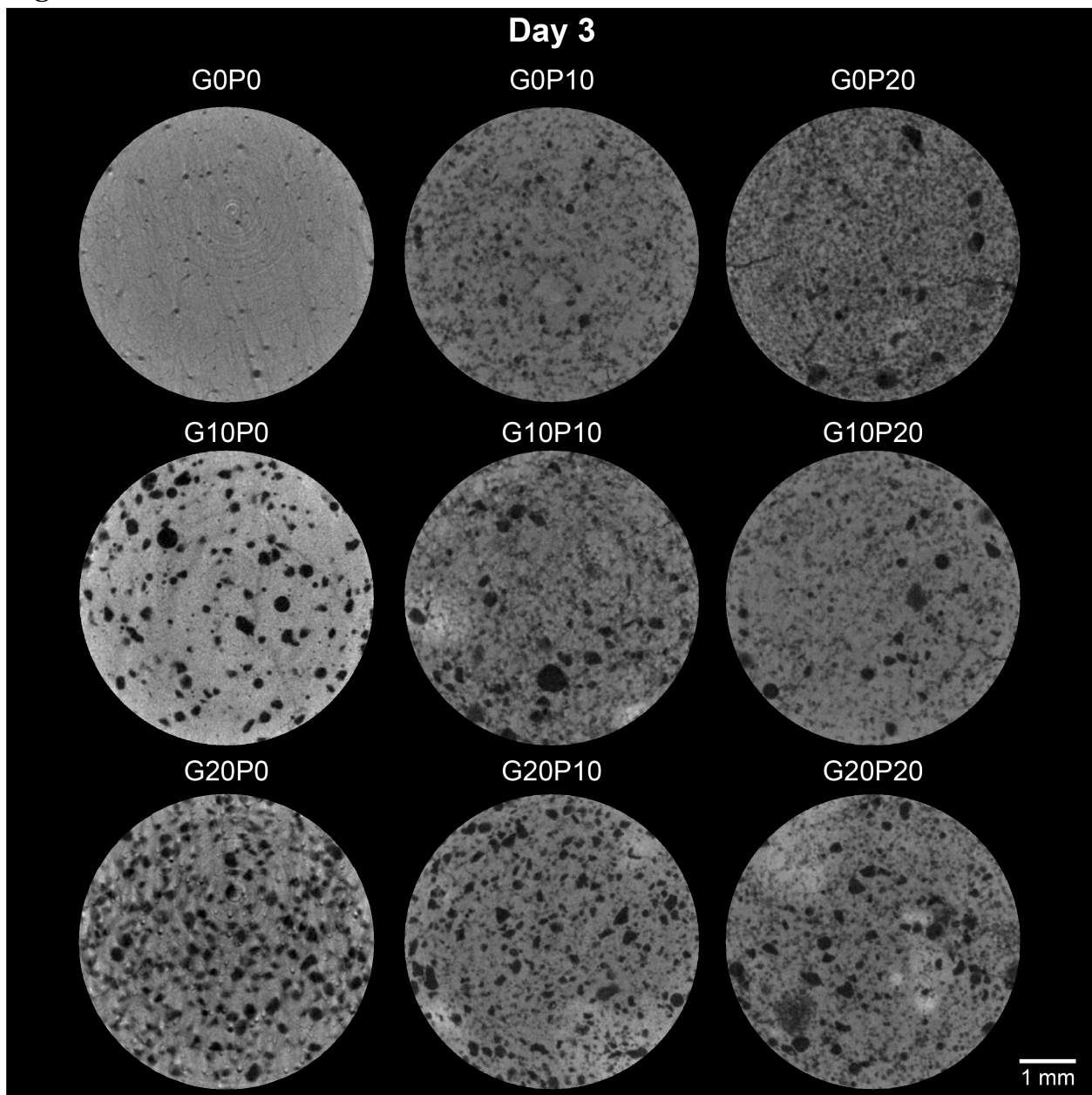
**Figure S2.** The molecular weight ( $M_w$ ) of the PLGA polymer inside CPC scaffolds evaluated by advanced polymer chromatography (APC) after fabrication (day 0) and after being leached in PBS at 37°C for 3 days, 4 wks, and 8 wks. GMPs and PLGA MPs were introduced in three different weight fractions (0, 10, and 20%). Those that do not share the same letter are statistically significantly different ( $p < 0.05$ ). At 8 wks, GOP10 and G10P10 (marked with an \*) could not be quantified due to chromatogram extending beyond that of the lowest standard (266 Da).

**Figure S3**

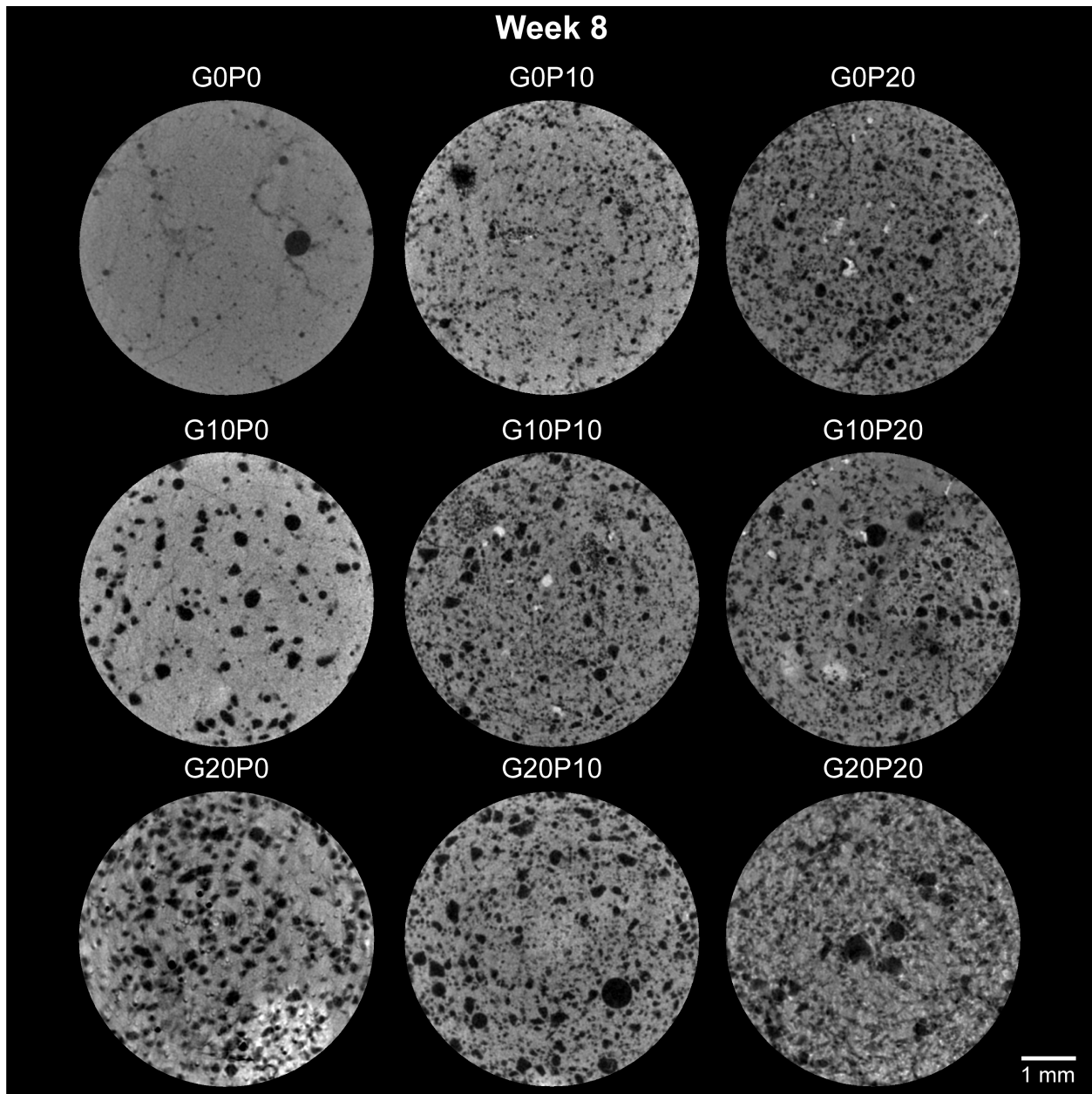


**Figure S3.** Representative scanning electron microscopy (SEM) images showing the internal morphology of composites upon fabrication (Day 0). The smooth spherical particles annotated by a yellow arrow represent PLGA MPs, while the jagged irregular particles marked with a green arrow represent GMPs. The white scale bar in the image represents 50  $\mu\text{m}$ .

*Figure S4 A*

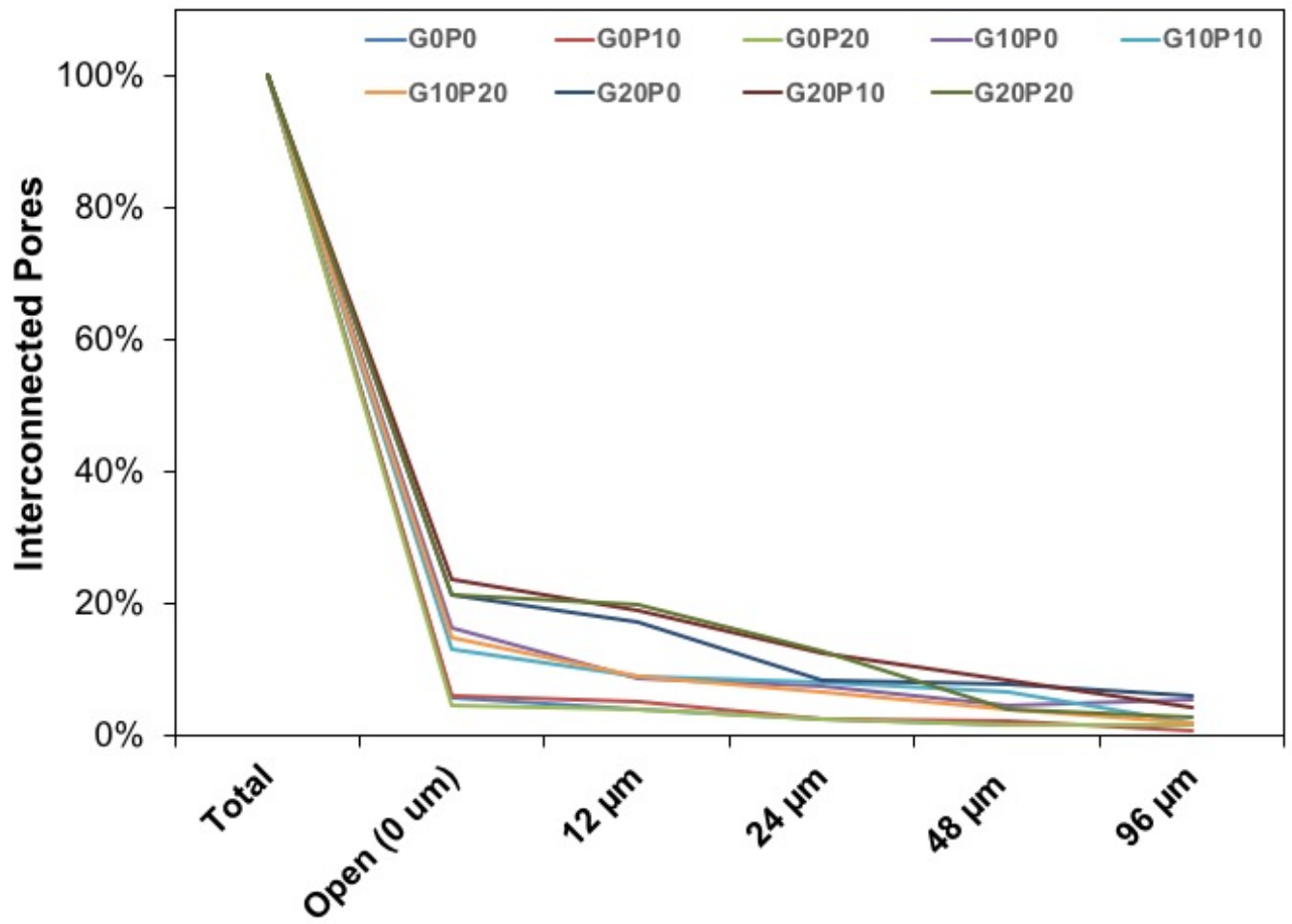


**Figure S4 B**

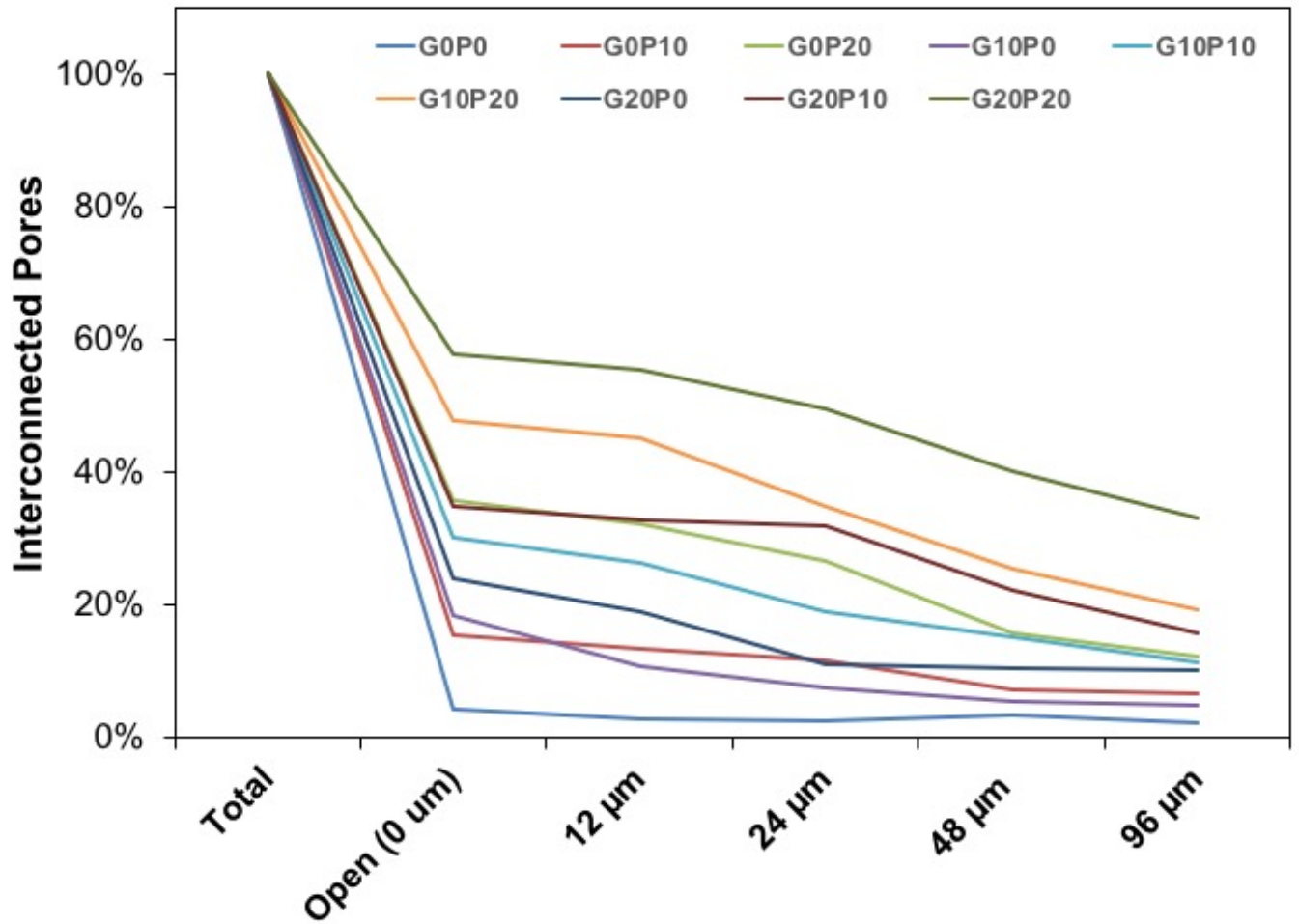


**Figure S4.** Representative  $\mu$ CT cross sections of GMP/PLGA/CPC composites after being leached for 3 days (A) and 8 weeks (B). Scale bar represents 1 mm.

Figure S5 A



**Figure S5 B**



**Figure S5.** Representative pore interconnectivity of the GMP/PLGA/CPC composites after being leached for 3 days (A) and 8 wks (B). In both graphs, the percentage of porosity available to allow the penetration of an object of a given size is presented.

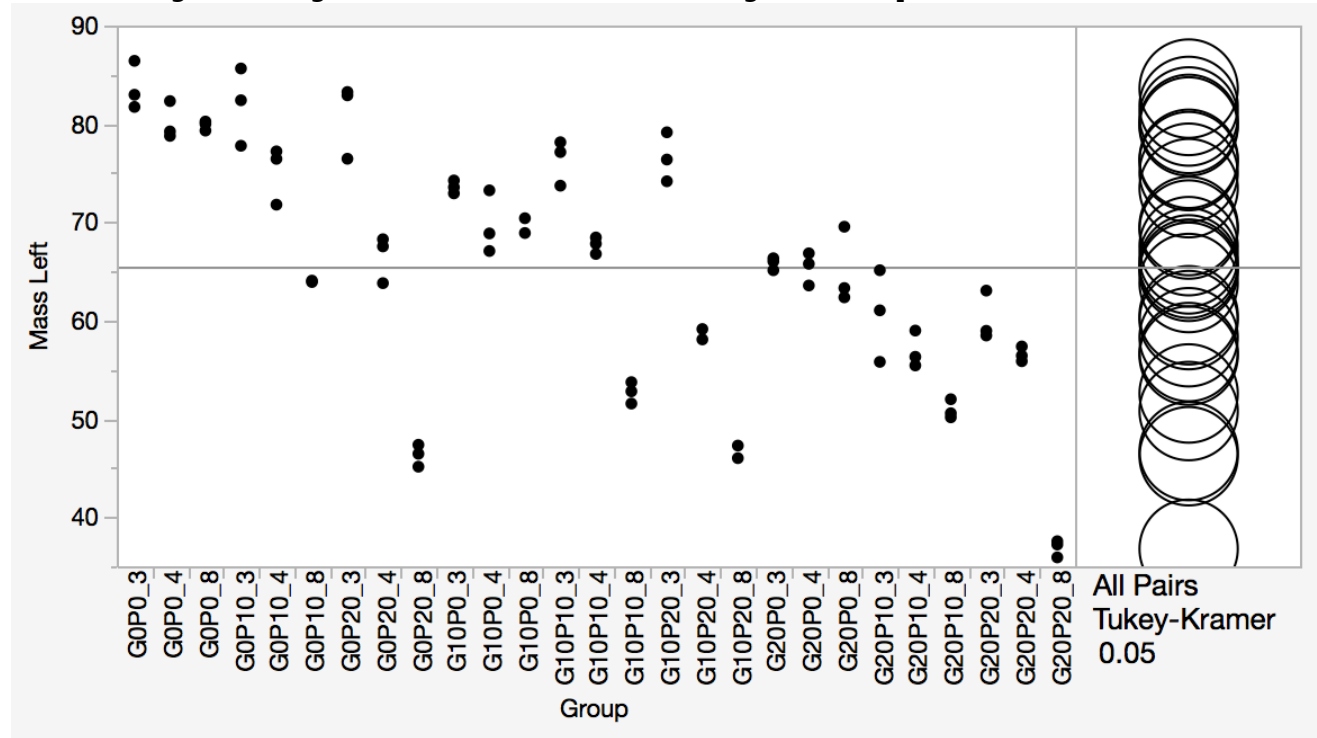


## Statistical Analysis

(Table S1) Mass Remaining Analysis (ANOVA)

### Fit Group

### Oneway Analysis of Mass Left By Group



### Means Comparisons

Comparisons for all pairs using Tukey-Kramer HSD

Confidence Quantile

q*	Alpha
3.89742	0.05

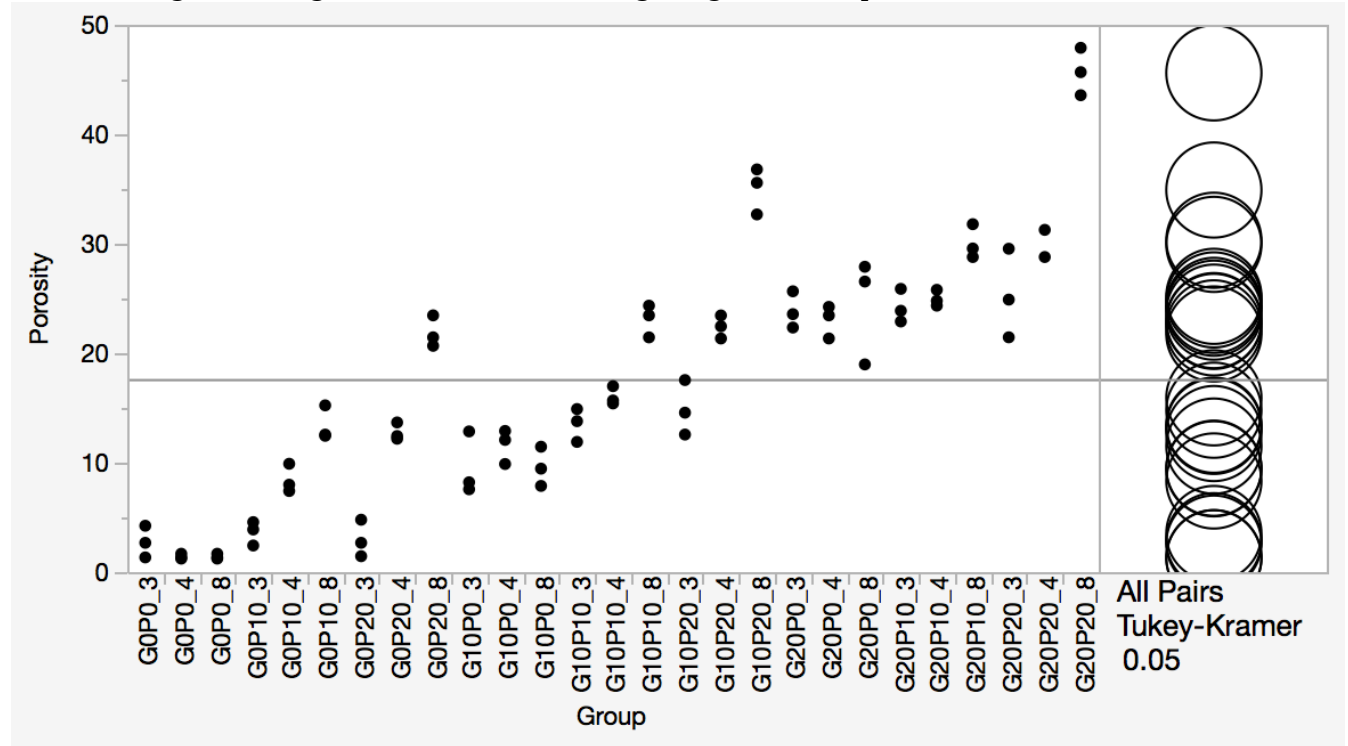
### Connecting Letters Report

Level		Mean
G0P0_3	A	83.696667
G0P10_3	A B	81.933333
G0P20_3	A B	80.863333
G0P0_4	A B C	80.106690
G0P0_8	A B C	79.846965
G10P20_3	B C D	76.556667
G10P10_3	B C D E	76.323333
G0P10_4	B C D E	75.140923
G10P0_3	C D E F	73.573333
G10P0_4	D E F G	69.718951
G10P0_8	E F G	69.361799
G10P10_4	F G H	67.674635
G0P20_4	F G H I	66.531196
G20P0_3	G H I	65.816667
G20P0_4	G H I J	65.383789
G20P0_8	G H I J	65.062973
G0P10_8	G H I J K	63.923890
G20P10_3	H I J K L	60.643333
G20P20_3	I J K L	60.153333
G10P20_4	J K L M	58.410180
G20P10_4	K L M N	56.901966
G20P20_4	L M N	56.543057
G10P10_8	M N O	52.690039
G20P10_8	N O	50.901707
G10P20_8	O	46.831728
G0P20_8	O	46.309273
G20P20_8	P	36.886561

Levels not connected by same letter are significantly different.

(Table S2) Porosity Analysis (ANOVA)

### Oneway Analysis of Porosity By Group



### Means Comparisons

Comparisons for all pairs using Tukey-Kramer HSD  
Confidence Quantile

q*	Alpha
3.89742	0.05

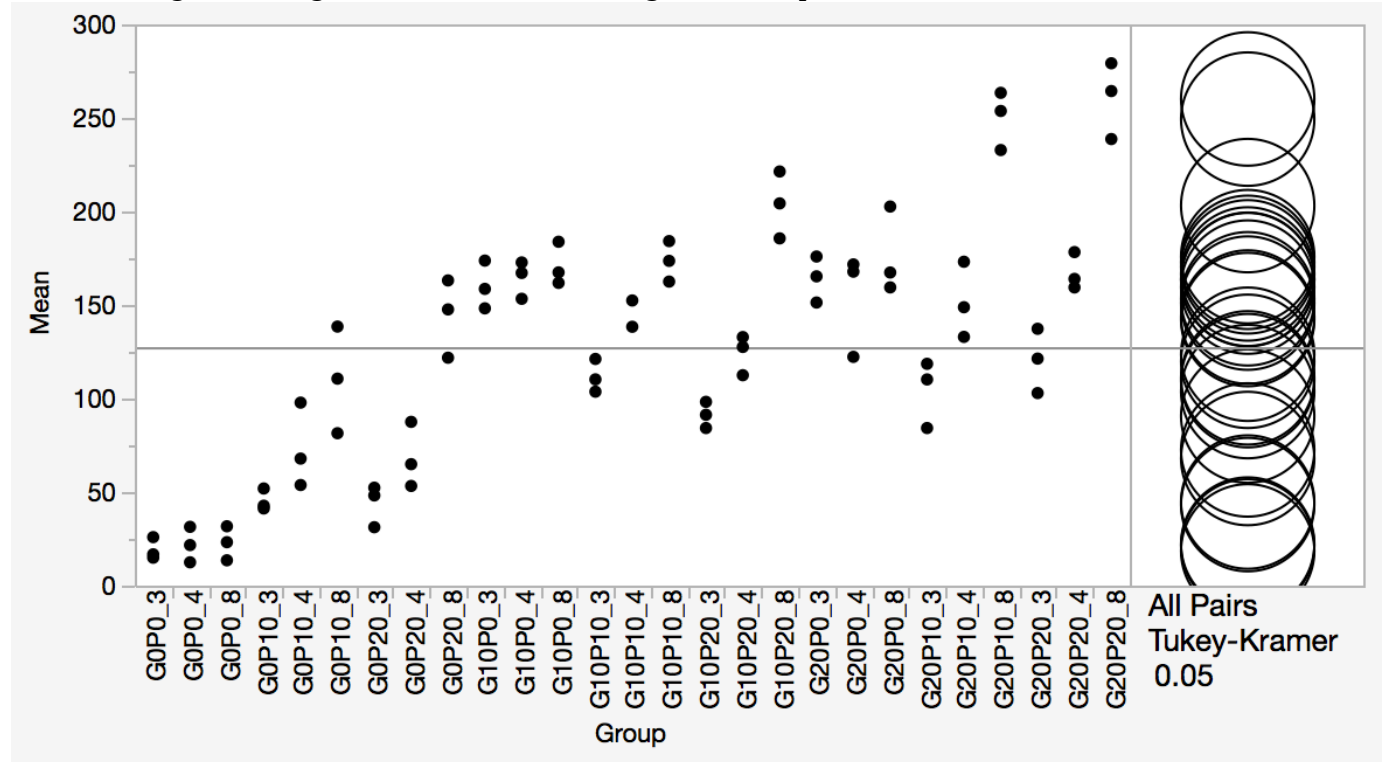
## Connecting Letters Report

Level		Mean
G20P20_8	A	45.713000
G10P20_8	B	35.011333
G20P20_4	B C	30.419933
G20P10_8	B C D	30.045567
G20P20_3	C D E	25.296667
G20P10_4	C D E	24.973189
G20P0_8	C D E	24.473000
G20P10_3	D E	24.216333
G20P0_3	E	23.855467
G10P10_8	E	23.081000
G20P0_4	E	23.009756
G10P20_4	E	22.416244
G0P20_8	E F	21.860533
G10P10_4	F G	16.025233
G10P20_3	G H	14.893600
G10P10_3	G H I	13.527867
G0P10_8	G H I	13.417100
G0P20_4	G H I	12.764244
G10P0_4	G H I	11.618156
G10P0_8	H I J	9.594833
G10P0_3	H I J	9.543693
G0P10_4	I J K	8.431644
G0P10_3	J K L	3.631233
G0P20_3	K L	2.973200
G0P0_3	K L	2.753377
G0P0_4	L	1.420053
G0P0_8	L	1.420053

Levels not connected by same letter are significantly different.

(Table S3) Mean Pore Size Analysis (ANOVA)

### Oneway Analysis of Mean By Group



### Means Comparisons

Comparisons for all pairs using Tukey-Kramer HSD  
Confidence Quantile

q*	Alpha
3.89742	0.05

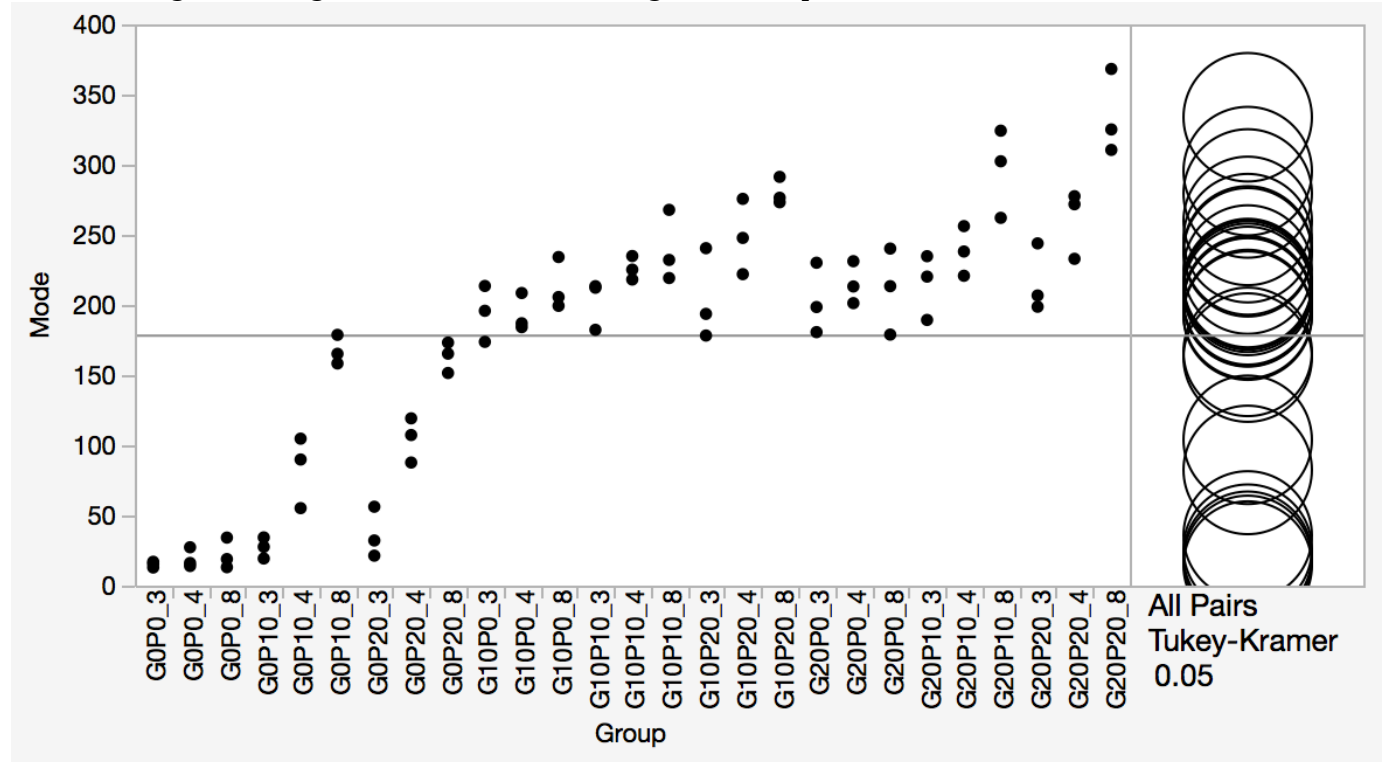
### Connecting Letters Report

Level	Mean
G20P20_8 A	260.76000
G20P10_8 A B	249.98667
G10P20_8 B C	203.76000
G20P0_8 C D	176.46667
G10P10_8 C D E	173.42000
G10P0_8 C D E	171.01667
G20P20_4 C D E F	167.20333
G10P0_4 C D E F	164.39333
G20P0_3 C D E F	164.18000
G10P0_3 C D E F G	160.18333
G20P0_4 C D E F G H	153.93000
G20P10_4 D E F G H	151.63000
G0P20_8 D E F G H	144.19667
G10P10_4 D E F G H	142.83667
G10P20_4 E F G H I	124.29667
G20P20_3 F G H I J	120.51333
G10P10_3 G H I J K	111.68667
G0P10_8 G H I J K	110.18333
G20P10_3 H I J K	104.33333
G10P20_3 I J K L	91.25667
G0P10_4 J K L M	73.15333
G0P20_4 K L M N	68.57000
G0P10_3 L M N	45.28667
G0P20_3 L M N	43.92000
G0P0_8 M N	22.82000
G0P0_4 N	21.84333
G0P0_3 N	19.18333

Levels not connected by same letter are significantly different.

(Table S4) Mode Pore Size Analysis (ANOVA)

### Oneway Analysis of Mode By Group



### Means Comparisons

Comparisons for all pairs using Tukey-Kramer HSD  
Confidence Quantile

q*	Alpha
3.89742	0.05

## Connecting Letters Report

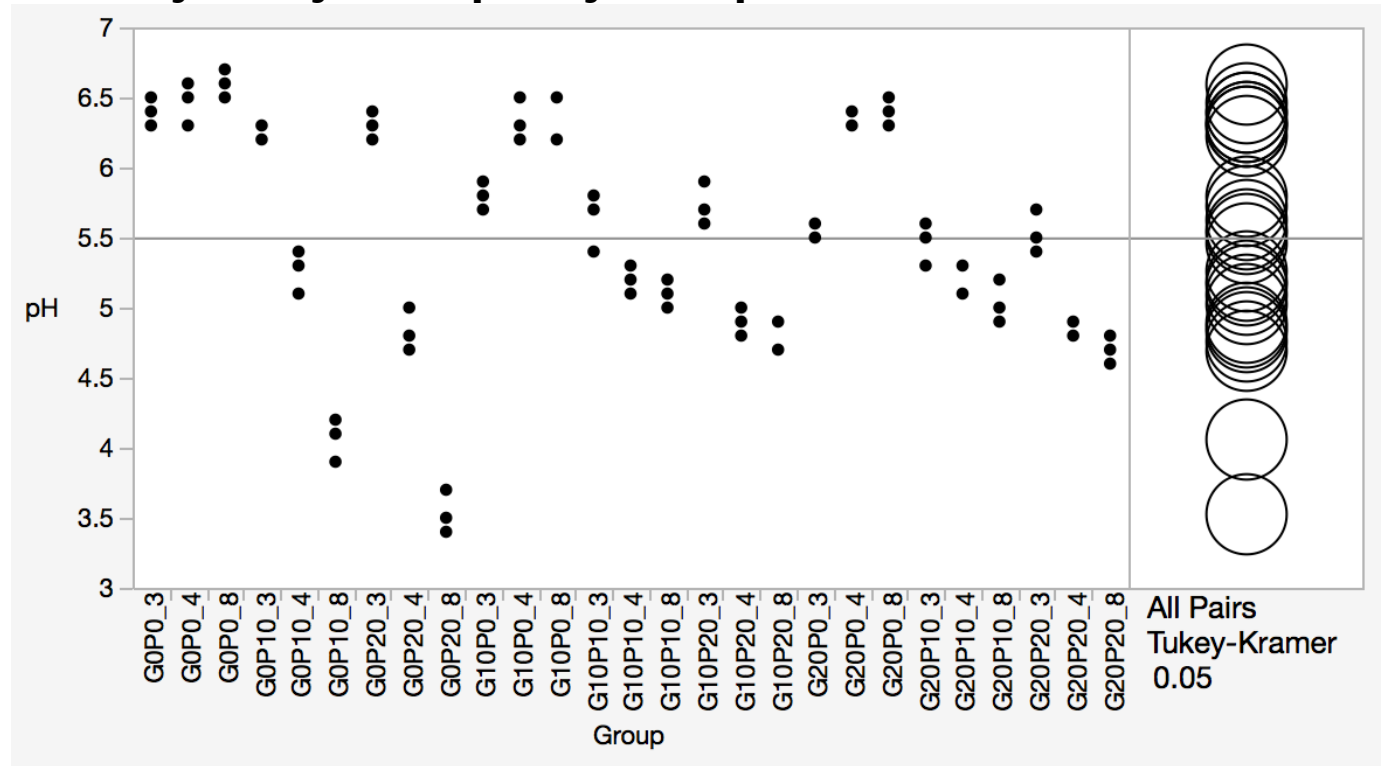
Level	Mean
G20P20_8 A	334.63000
G20P10_8 A B	296.26000
G10P20_8 A B C	280.31667
G20P20_4 B C D	260.75333
G10P20_4 B C D E	248.47667
G10P10_8 B C D E	239.71333
G20P10_4 B C D E	238.38667
G10P10_4 C D E F	226.09333
G20P20_3 C D E F	216.48333
G20P0_4 D E F	215.24667
G20P10_3 D E F	214.78333
G10P0_8 D E F	213.08667
G20P0_8 D E F	210.84333
G10P20_3 D E F	204.13000
G20P0_3 D E F	203.11000
G10P10_3 D E F	202.67667
G10P0_3 E F	194.39000
G10P0_4 E F	193.24333
G0P10_8 F G	167.42000
G0P20_8 F G	163.35333
G0P20_4 G H	104.75000
G0P10_4 H I	83.27000
G0P20_3 I J	36.57000
G0P10_3 I J	27.11000
G0P0_8 I J	22.08000
G0P0_4 I J	19.14000
G0P0_3 J	15.11000

Levels not connected by same letter are significantly different.



*(Table S5) Incubation Media pH Analysis (ANOVA)*

### Oneway Analysis of pH By Group



### Means Comparisons

Comparisons for all pairs using Tukey-Kramer HSD

Confidence Quantile

q*	Alpha
3.89742	0.05

## Connecting Letters Report

Level	Mean
G0P0_8 A	6.6000000
G0P0_4 A	6.4666667
G0P0_3 A	6.4000000
G20P0_8 A	6.4000000
G10P0_4 A	6.3333333
G20P0_4 A	6.3333333
G0P20_3 A	6.3000000
G10P0_8 A	6.3000000
G0P10_3 A	6.2333333
G10P0_3 B	5.8000000
G10P20_3 B	5.7333333
G10P10_3 B C	5.6333333
G20P0_3 B C D	5.5666667
G20P20_3 B C D	5.5333333
G20P10_3 B C D E	5.4666667
G0P10_4 C D E F	5.2666667
G10P10_4 D E F G	5.2000000
G20P10_4 D E F G H	5.1666667
G10P10_8 E F G H I	5.1000000
G20P10_8 F G H I	5.0333333
G10P20_4 F G H I	4.9000000
G20P20_4 F G H I	4.8666667
G0P20_4 G H I	4.8333333
G10P20_8 H I	4.7666667
G20P20_8 I	4.7000000
G0P10_8 J	4.0666667
G0P20_8 K	3.5333333

Levels not connected by same letter are significantly different.