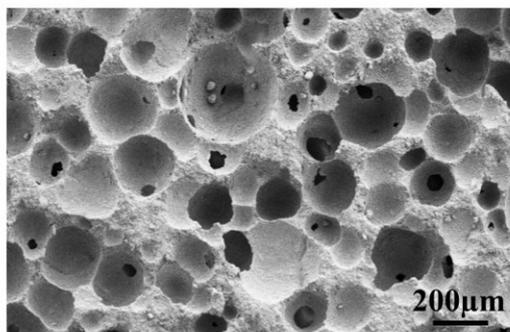


Fig. S1 XRD patterns of BCP ceramics.

A



B

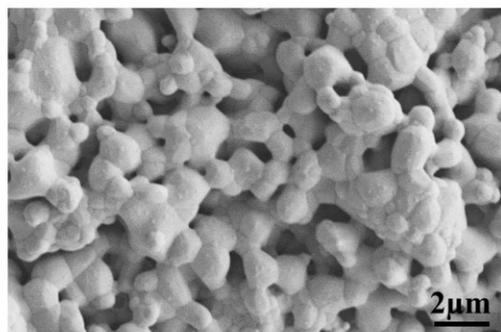


Fig. S2 SEM images of BCP ceramics.

Table S1. Inflammatory genes expressed (H value) by RAW 264.7 macrophages and the F value under different MVs parameters. * $p < 0.05$, ** $p < 0.01$ represents significant difference among three levels for one factor.

Group	H value					
	D3			D5		
	<i>Il1β</i>	<i>Tnfα</i>	<i>Il10</i>	<i>Il1β</i>	<i>Tnfα</i>	<i>Il10</i>
1	2.8205	2.3615	2.5106	2.5022	2.4207	2.8727
2	2.1591	1.9203	2.4990	2.7083	3.8582	3.5646
3	3.2832	1.8550	3.8705	2.1546	2.8443	3.1717
4	3.1747	2.5959	2.1818	2.7658	3.2486	4.0208
5	4.4455	2.5992	3.8997	2.7730	3.1380	4.0865
6	3.0014	3.2861	2.6142	2.7964	2.3857	3.9431
7	2.6283	3.5864	2.8110	2.8503	3.6730	4.2747
8	2.8952	2.7114	2.7612	4.1786	3.9354	3.2688
9	2.5885	2.0610	2.9697	3.2624	2.9572	3.9717
Factor	F value					
A (Frequency)	6.8474**	11.4052**	0.1055	11.6070**	11.2449**	8.3506**
B (Magnitude)	0.7045	3.8471*	4.0711*	3.8981*	23.6349**	0.0812
C (Time)	5.3286*	5.9131*	9.7579**	3.7915*	5.7059*	3.6064

Table S2. Weight matrix (ω value) of inflammatory genes for RAW 264.7 cells under different MVs parameters.

Level	D3			D5			Average of weight matrix
	<i>Il1β</i>	<i>Tnfα</i>	<i>Il10</i>	<i>Il1β</i>	<i>Tnfα</i>	<i>Il10</i>	
A ₁	0.1541	0.1744	0.0220	0.1825	0.1054	0.1698	0.1572
A ₂	0.1199	0.1262	0.0215	0.1613	0.1096	0.2130	0.1460
A ₃	0.1570	0.1280	0.0211	0.1306	0.0910	0.2035	0.1420
B ₁	0.0522	0.0728	0.1073	0.0883	0.1563	0.0200	0.1062
B ₂	0.0474	0.0860	0.1310	0.0742	0.1336	0.0196	0.1062
B ₃	0.0508	0.0864	0.1353	0.0873	0.1783	0.0199	0.1218
C ₁	0.1421	0.0984	0.1695	0.0834	0.0813	0.1075	0.1191
C ₂	0.1564	0.1250	0.1644	0.0904	0.0706	0.1232	0.1205
C ₃	0.1196	0.1023	0.2274	0.1016	0.0736	0.1229	0.1306

Table S3. Osteogenic marker expressed (H value) by BMMSCs and the F value under different CM derived from RAW 264.7 cells response to different MVs parameters. * $p < 0.05$, ** $p < 0.01$ represents significant difference among three levels for one factor.

Group	H value					
	<i>Alp</i> protein	<i>Runx2</i>	<i>Col I</i>	<i>Alp</i>	<i>Opn</i>	<i>Osx</i>
1	28.1706	3.1544	3.8214	3.2886	2.9602	2.5611
2	40.3267	2.6181	2.8453	2.2539	1.9324	1.4393
3	31.7241	2.5614	2.7951	3.2908	3.0058	2.6566
4	37.7220	2.4423	2.8162	2.5792	3.0169	2.3189
5	42.9139	2.3912	2.7450	1.8681	2.3915	1.4386
6	53.1790	2.6206	2.8176	1.9173	2.3951	1.7608
7	32.8309	3.2623	3.2266	1.8237	2.4153	1.5108
8	22.9085	2.2797	2.9162	1.3865	2.4123	1.4294
9	22.2112	2.0457	3.8535	1.4598	2.2954	1.3603
Factor	F value					
A (Frequency)	24.7331**	1.8223	5.4794*	17.2937**	0.9906	16.2401**
B (Magnitude)	0.6592	6.9312**	3.9296*	4.70636*	3.8291*	13.4064**
C (Time)	0.4079	2.9127	1.5911	0.4719	0.5518	1.28430

Table S4. Weight matrix (ω value) of osteogenic markers for BMSCs under different CM derived from RAW 264.7 cells response to different MVs parameters.

Level	Alp protein	<i>Runx2</i>	<i>Col I</i>	<i>Alp</i>	<i>Opn</i>	<i>Osx</i>	Average of weight matrix
A ₁	0.2511	0.0866	0.1960	0.2630	0.0894	0.1876	0.2244
A ₂	0.3352	0.0774	0.1736	0.1895	0.0883	0.1555	0.2135
A ₃	0.1953	0.0788	0.2071	0.1390	0.0806	0.1212	0.1657
B ₁	0.0371	0.1707	0.0502	0.1201	0.2030	0.1593	0.1407
B ₂	0.0399	0.1405	0.0433	0.0860	0.1629	0.1073	0.1080
B ₃	0.0403	0.1393	0.0482	0.1041	0.1861	0.1440	0.1243
C ₁	0.0337	0.1055	0.0965	0.0325	0.0644	0.0435	0.0627
C ₂	0.0324	0.0931	0.0961	0.0310	0.0600	0.0387	0.0585
C ₃	0.0347	0.1076	0.0885	0.0344	0.0648	0.0424	0.0621

The index layer matrix: M , factor layer matrix: T , horizontal layer matrix: S , and the weight value ω of all experimental results are listed as followed:

ELISA-II-1 β -D3

```
M=np.array([[0.007958223,0,0],[0.008148684,0,0],[0.007349202,0,0],[0,0.008060993,0],[0,0.007891153,0],[0,0.007481323,0],[0,0,0.007874561],[0,0,0.007587415],[0,0,0.007958223]])
```

```
T=np.array([[42.63281799,0,0],[0,42.67400765,0],[0,0,42.69818634]])
```

```
S=np.array([[0.169230769,0,0],[0,0.553846154,0],[0,0,0.276923077]])
```

```
 $\omega$ =array([[0.05741686, 0.        ],  
        [0.058791, 0.        ],  
        [0.05302291, 0.        ],  
        [0.        , 0.19052024, 0.        ],  
        [0.        , 0.1865061, 0.        ],  
        [0.        , 0.17681983, 0.        ],  
        [0.        , 0.        , 0.0931097 ],  
        [0.        , 0.        , 0.08971445],  
        [0.        , 0.        , 0.09409893]])
```

ELISA-II-1 β -D5

```
M=np.array([[0.007249623,0,0],[0.008461654,0,0],[0.007696559,0,0],[0,0.007958223,0],[0,0.007874561,0],[0,0.007496297,0],[0,0,0.00811338],[0,0,0.007602817],[0,0,0.007618281]])
```

```
T=np.array([[42.72073628,0,0],[0,42.86495518,0],[0,0,42.85504099]])
```

```
S=np.array([[0.552238806,0,0],[0,0.21641791,0],[0,0,0.231343284]])
```

```
 $\omega$ =array([[0.17103346, 0.        ],  
        [0.19962775, 0.        ],  
        [0.18157759, 0.        ],  
        [0.        , 0.0738264, 0.        ],  
        [0.        , 0.07305029, 0.        ],  
        [0.        , 0.06954123, 0.        ],  
        [0.        , 0.        , 0.08043788],  
        [0.        , 0.        , 0.07537605],  
        [0.        , 0.        , 0.07552936]])
```

ELISA-Tnf- α -D3

```
M=np.array([[0.000286426,0,0],[0.00027771,0,0],[0.000271174,0,0],[0,0.000282271,0],[0,0.000276555,0],[0,0.00276148,0],[0,0,0.000282078],[0,0,0.000285356],[0,0,0.000268072]])
```

```
T=np.array([[3591.482753,0,0],[0,1197.641845,0],[0,0,1196.879742]])
```

```
S=np.array([[0.392061955,0,0],[0,0.156824782,0],[0,0,0.451113262]])
```

```
 $\omega$ =array([[0.4033118, 0.        ],  
        [0.39103894, 0.        ],  
        [0.3818357, 0.        ],  
        [0.        , 0.05301612, 0.        ],  
        [0.        , 0.05194254, 0.        ],  
        [0.        , 0.0518661, 0.        ],
```

```
[0.      , 0.      , 0.1523019 ],
[0.      , 0.      , 0.15407179],
[0.      , 0.      , 0.14473967]]])
```

ELISA-Tnf- α -D5

```
M=np.array([[0.00023728,0,0],[0.000236518,0,0],[0.000221363,0,0],[0,0.000231241,0],[0,0.000226194,0],[0,0.0023728,0],[0,0,0.000222918],[0,0,0.000234368],[0,0,0.00023769]])
T=np.array([[1438.514285,0,0],[0,1439.437655,0],[0,0,1438.896773]])
S=np.array([[0.384378844,0,0],[0,0.26199262,0],[0,0,0.353628536]])
 $\omega$ =array([[0.13120029, 0.      , 0.      ],
          [0.13077895, 0.      , 0.      ],
          [0.12239923, 0.      , 0.      ],
          [0.      , 0.08720608, 0.      ],
          [0.      , 0.08530274, 0.      ],
          [0.      , 0.08948352, 0.      ],
          [0.      , 0.      , 0.11342847],
          [0.      , 0.      , 0.11925463],
          [0.      , 0.      , 0.12094498]])
```

PCR-*IL1 β* -D3

```
M=np.array([[0.121022854,0,0],[0.094146373,0,0],[0.12327279,0,0],[0,0.115960254,0],[0,0.1052641,0],[0,0.112699026,0],[0,0,0.114715686],[0,0,0.126224394],[0,0,0.096551676]])
T=np.array([[2.954715878,0,0],[0,2.994698961,0],[0,0,2.963035334]])
S=np.array([[0.431163464,0,0],[0,0.150544172,0],[0,0,0.418292364]])
 $\omega$ =array([[0.15417894, 0.      , 0.      ],
          [0.11993923, 0.      , 0.      ],
          [0.15704529, 0.      , 0.      ],
          [0.      , 0.05227888, 0.      ],
          [0.      , 0.04745669, 0.      ],
          [0.      , 0.05080861, 0.      ],
          [0.      , 0.      , 0.14218035],
          [0.      , 0.      , 0.15644441],
          [0.      , 0.      , 0.1196676 ]])
```

PCR-Tnf α -D3

```
M=np.array([[0.162949006,0,0],[0.117906053,0,0],[0.119630611,0,0],[0,0.117041726,0],[0,0.138292733,0],[0,0.138844862,0],[0,0,0.119629093],[0,0,0.152036373],[0,0,0.124366686]])
T=np.array([[2.496968244,0,0],[0,2.536916445,0],[0,0,2.525047512]])
S=np.array([[0.428762965,0,0],[0,0.245372942,0],[0,0,0.325864093]])
 $\omega$ =array([[0.17445443, 0.      , 0.      ],
          [0.12623111, 0.      , 0.      ],
          [0.12807743, 0.      , 0.      ],
          [0.      , 0.07285738, 0.      ],
          [0.      , 0.08608593, 0.      ],
```

```
[0.          , 0.08642963, 0.          ],
[0.          , 0.          , 0.09843349],
[0.          , 0.          , 0.12509892],
[0.          , 0.          , 0.10233168]])
```

PCR-*IL10*-D3

```
M=np.array([[8.880279847,0,0],[8.695743644,0,0],[8.542100846,0,0],[0,7.503542571,0],[0,9.160000628,0],[0,9.454581138,0],[0,0,7.886129667],[0,0,7.650655969],[0,0,10.5813387]])
T=np.array([[0.038287589,0,0],[0,0.038287589,0],[0,0,0.038287589]])
S=np.array([[0.064786487,0,0],[0,0.373769316,0],[0,0,0.561444197]])
ω=array([[0.0220277 , 0.          , 0.          ],
         [0.02156995, 0.          , 0.          ],
         [0.02118884, 0.          , 0.          ],
         [0.          , 0.10738114, 0.          ],
         [0.          , 0.13108626, 0.          ],
         [0.          , 0.13530192, 0.          ],
         [0.          , 0.          , 0.16952296],
         [0.          , 0.          , 0.16446114],
         [0.          , 0.          , 0.2274601 ]])
```

PCR-*IL1β*-D5

```
M=np.array([[0.135773174,0,0],[0.119972097,0,0],[0.097167691,0,0],[0,0.123176503,0],[0,0.10352043,0],[0,0.121748956,0],[0,0,0.105515355],[0,0,0.114461091],[0,0,0.128566161]])
T=np.array([[2.833559846,0,0],[0,2.869886072,0],[0,0,2.869089687]])
S=np.array([[0.47450662,0,0],[0,0.24996083,0],[0,0,0.27553255]])
ω=array([[0.18255286, 0.          , 0.          ],
         [0.16130763, 0.          , 0.          ],
         [0.13064613, 0.          , 0.          ],
         [0.          , 0.08836179, 0.          ],
         [0.          , 0.07426132, 0.          ],
         [0.          , 0.08733772, 0.          ],
         [0.          , 0.          , 0.0834128 ],
         [0.          , 0.          , 0.09048465],
         [0.          , 0.          , 0.1016351 ]])
```

PCR-*Tnfα*-D5

```
M=np.array([[0.109609654,0,0],[0.11399371,0,0],[0.094645613,0,0],[0,0.107038251,0],[0,0.091477427,0],[0,0.122140169,0],[0,0,0.11439201],[0,0,0.099362397],[0,0,0.103569103]])
T=np.array([[3.142193917,0,0],[0,3.118608348,0],[0,0,3.151358051]])
S=np.array([[0.306028742,0,0],[0,0.468321241,0],[0,0,0.225650016]])
ω=array([[0.10540082, 0.          , 0.          ],
         [0.10961654, 0.          , 0.          ],
         [0.09101138, 0.          , 0.          ],
         [0.          , 0.15633049, 0.          ],
```

```

[0.      , 0.13360375, 0.      ],
[0.      , 0.178387  , 0.      ],
[0.      , 0.      , 0.08134462],
[0.      , 0.      , 0.070657  ],
[0.      , 0.      , 0.0736484 ]])

```

PCR-//10-D5

```

M=np.array([[9.60913429,0,0],[12.05054257,0,0],[11.51539213,0,0],[0,11.16838539,0],[0,10.91998859,0],[0,11.08669501,0],[0,0,10.08466329],[0,0,11.55729697],[0,0,11.53310872]])
T=np.array([[0.030143117,0,0],[0,0.030143117,0],[0,0,0.030143117]])
S=np.array([[0.58653314,0,0],[0,0.059675786,0],[0,0,0.3537910743]])
ω=array([[0.16988889, 0.      , 0.      ],
        [0.21305283, 0.      , 0.      ],
        [0.20359141, 0.      , 0.      ],
        [0.      , 0.02008985, 0.      ],
        [0.      , 0.01964303, 0.      ],
        [0.      , 0.0199429 , 0.      ],
        [0.      , 0.      , 0.10754654],
        [0.      , 0.      , 0.12325124],
        [0.      , 0.      , 0.12299329]])

```

Protein-Alp- D7

```

M=np.array([[100.2215871,0,0],[133.8149602,0,0],[77.95076562,0,0],[0,98.72364553,0],[0,106.1492169,0],[0,107.1144504,0],[0,0,104.2582068],[0,0,100.260021],[0,0,107.4690852]])
T=np.array([[0.003205259,0,0],[0,0.003205259,0],[0,0,0.003205259]])
S=np.array([[0.781710299,0,0],[0,0.117412927,0],[0,0,0.100876774]])
ω=array([[0.2511136 , 0.      , 0.      ],
        [0.33528462, 0.      , 0.      ],
        [0.19531219, 0.      , 0.      ],
        [0.      , 0.03715354, 0.      ],
        [0.      , 0.03994807, 0.      ],
        [0.      , 0.04031133, 0.      ],
        [0.      , 0.      , 0.03371045],
        [0.      , 0.      , 0.03241769],
        [0.      , 0.      , 0.03474864]])

```

PCR-Alp- D7

```

M=np.array([[8.833523968,0,0],[6.364733947,0,0],[4.670071119,0,0],[0,7.691635044,0],[0,5.508608185,0],[0,6.668085805,0],[0,0,6.592544959],[0,0,6.293029784],[0,0,6.982754292]])
T=np.array([[0.050331359,0,0],[0,0.050331359,0],[0,0,0.050331359]])
S=np.array([[0.591718592,0,0],[0,0.310256324,0],[0,0,0.098025084]])
ω=array([[0.26308002, 0.      , 0.      ],
        [0.18955451, 0.      , 0.      ],
        [0.13908406, 0.      , 0.      ],

```

```

[0.      , 0.12010967, 0.      ],
[0.      , 0.08602035, 0.      ],
[0.      , 0.10412631, 0.      ],
[0.      , 0.      , 0.03252587],
[0.      , 0.      , 0.03104815],
[0.      , 0.      , 0.03445106]])

```

PCR-Col-I- D7

```

M=np.array([[9.461954351,0,0],[8.378901355,0,0],[9.996496329,0,0],[0,9.864361534,0],[0,8.506667502,0],[0,9.466322999,0],[0,0,9.55534728],[0,0,9.515179615],[0,0,8.76682514]])
T=np.array([[0.035922957,0,0],[0,0.035922957,0],[0,0,0.035922957]])
S=np.array([[0.576856343,0,0],[0,0.141945949,0],[0,0,0.281197707]])
omega=array([[0.19607427, 0.      , 0.      ],
              [0.17363082, 0.      , 0.      ],
              [0.20715125, 0.      , 0.      ],
              [0.      , 0.05029955, 0.      ],
              [0.      , 0.0433765 , 0.      ],
              [0.      , 0.0482699 , 0.      ],
              [0.      , 0.      , 0.09652289],
              [0.      , 0.      , 0.09611714],
              [0.      , 0.      , 0.08855767]])

```

PCR-Opn- D7

```

M=np.array([[7.898517036,0,0],[7.803614316,0,0],[7.123101018,0,0],[0,8.392441433,0],[0,6.736341729,0],[0,7.696449207,0],[0,0,7.767647407],[0,0,7.244823538],[0,0,7.812761424]])
T=np.array([[0.043811164,0,0],[0,0.043811164,0],[0,0,0.043811164]])
S=np.array([[0.25851909,0,0],[0,0.552133795,0],[0,0,0.189347115]])
omega=array([[0.08945878, 0.      , 0.      ],
              [0.08838391, 0.      , 0.      ],
              [0.0806764 , 0.      , 0.      ],
              [0.      , 0.20301 , 0.      ],
              [0.      , 0.16294958, 0.      ],
              [0.      , 0.18617421, 0.      ],
              [0.      , 0.      , 0.06443666],
              [0.      , 0.      , 0.06009956],
              [0.      , 0.      , 0.0648109 ]])

```

PCR-Osx- D7

```

M=np.array([[6.657140081,0,0],[5.518422778,0,0],[4.300651756,0,0],[0,6.390944469,0],[0,4.307465038,0],[0,5.777805108,0],[0,0,5.751398303],[0,0,5.118708492],[0,0,5.60610782]])
T=np.array([[0.060693553,0,0],[0,0.060693553,0],[0,0,0.060693553]])
S=np.array([[0.464547093,0,0],[0,0.410727396,0],[0,0,0.124725512]])
omega=array([[0.18769816, 0.      , 0.      ],
              [0.15559201, 0.      , 0.      ],

```

```
[0.12125693, 0.          , 0.          ],
[0.          , 0.15931669, 0.          ],
[0.          , 0.10737866, 0.          ],
[0.          , 0.14403204, 0.          ],
[0.          , 0.          , 0.04353828],
[0.          , 0.          , 0.0387488 ],
[0.          , 0.          , 0.04243843]])
```

PCR-Runx2- D7

```
M=np.array([[8.334029967,0,0],[7.454264806,0,0],[7.587892539,0,0],[0,8.85911499,0],[0,7.289205235,0],[0,7.27867087,0],[0,0,8.054921214],[0,0,7.106164751],[0,0,8.215101347]])
```

```
T=np.array([[0.042778576,0,0],[0,0.042778576,0],[0,0,0.042778576]])
```

```
S=np.array([[0.243032429,0,0],[0,0.450627234,0],[0,0,0.306340336]])
```

```
 $\omega$ =array([[0.08664542, 0.          , 0.          ],
[0.07749887, 0.          , 0.          ],
[0.07888814, 0.          , 0.          ],
[0.          , 0.17077886, 0.          ],
[0.          , 0.1405154 , 0.          ],
[0.          , 0.13933298, 0.          ],
[0.          , 0.          , 0.10555816],
[0.          , 0.          , 0.09312489],
[0.          , 0.          , 0.10765729]])
```