

Multifunctional 3D sponge-like macroporous cryogel-modified long carbon fiber reinforced polyetheretherketone implant with enhanced vascularization and osseointegration

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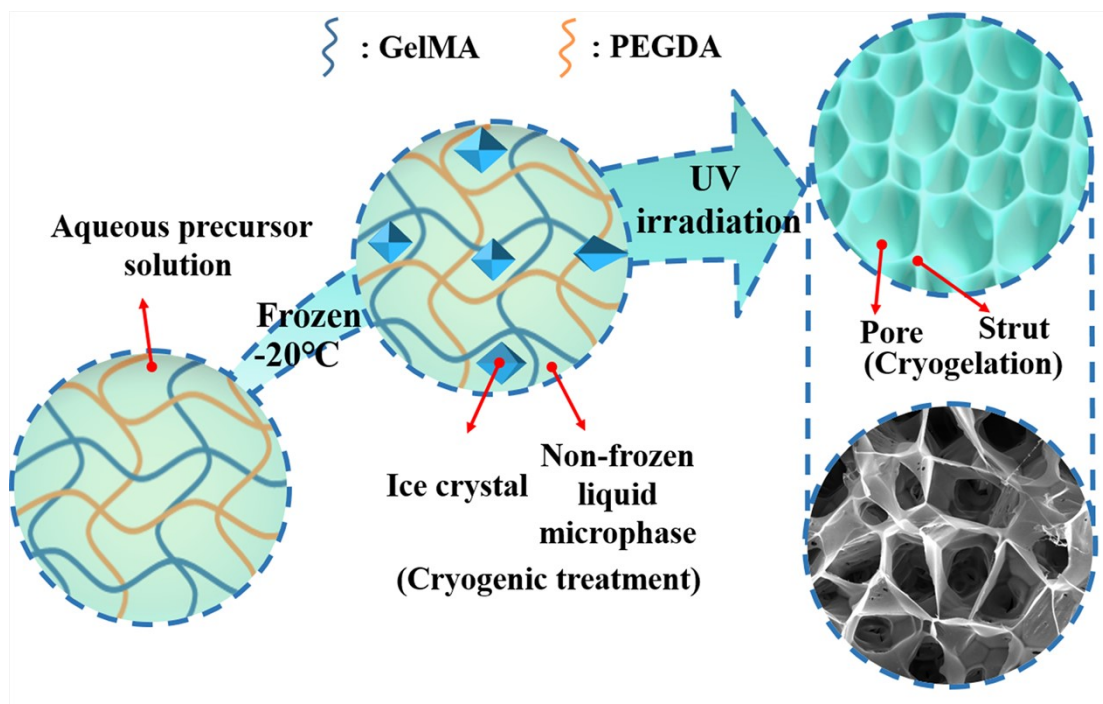


Fig. S1. Schematic depiction of the GelMA/PEGDA cryogel on the surface of the sulfonated LCFRPEEK formation.

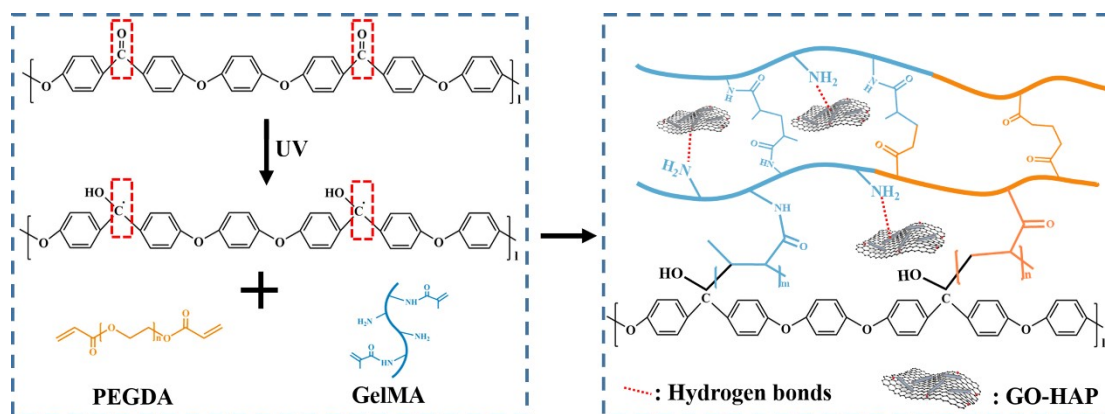


Fig. S2. Schematic illustration of the free radical polymerization on the surface of LCFRPEEK.

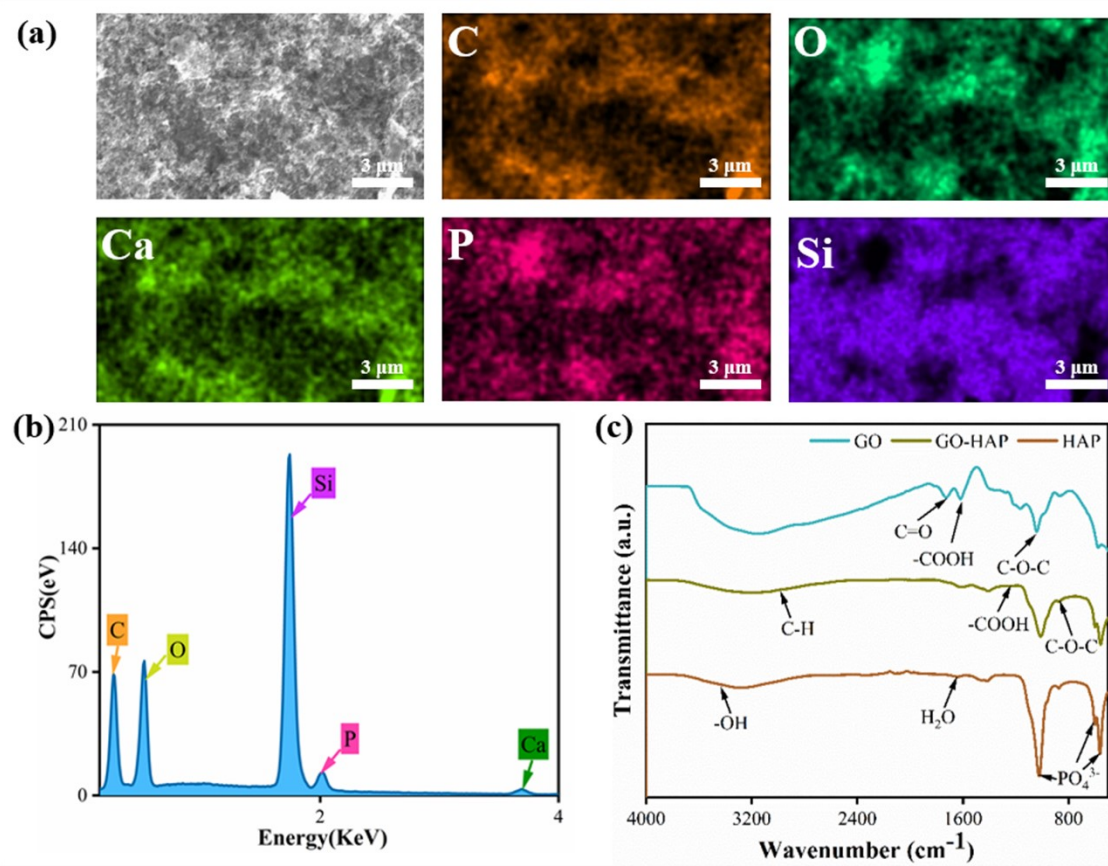


Fig. S3. (a) EDS images and (b) spectra of GO-HAP nanocomposite. (c) ATR-FTIR spectra of GO, HAP, and GO-HAP.

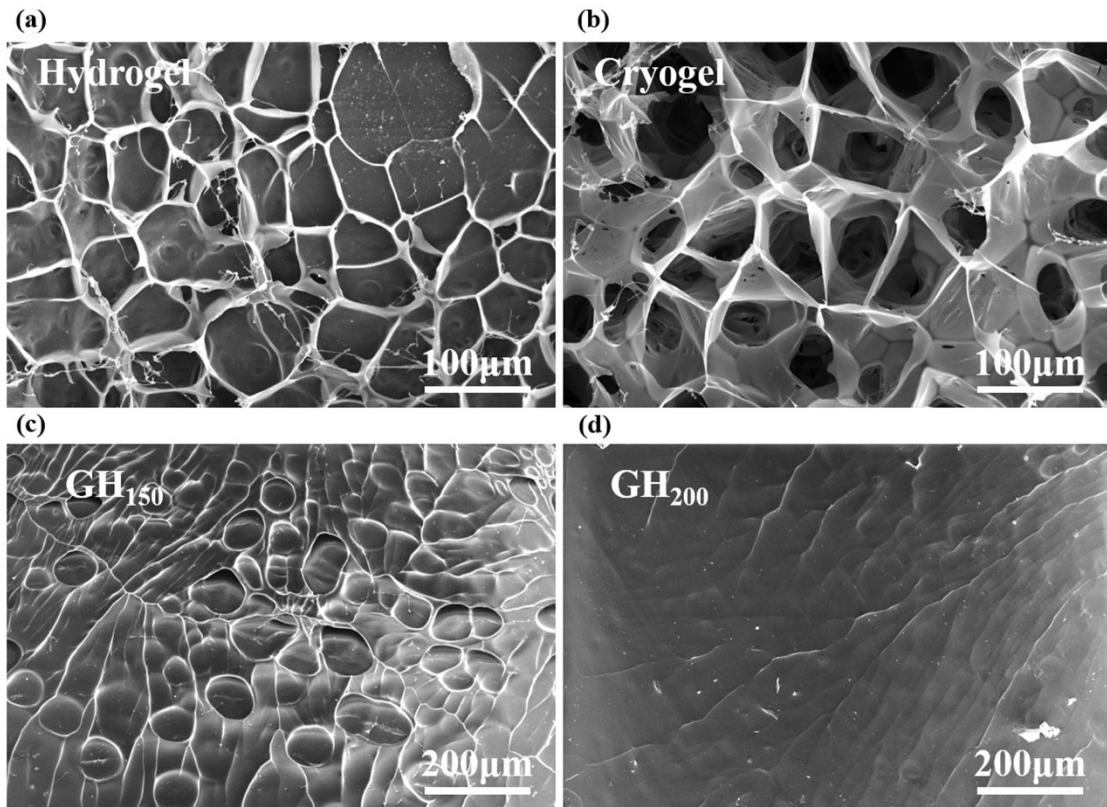


Fig. S4. (a) SEM images of GelMA/PEGDA hydrogel on the surface of sulfonated LCFRPEEK. [GelMA: PEGDA=10:5 wt%, the preparation method of GelMA/PEGDA hydrogel on the surface of sulfonated LCFRPEEK was similar to cryogel whereas without the freezing step.] (b) SEM images of GelMA/PEGDA cryogel on the surface of sulfonated LCFRPEEK. [GelMA: PEGDA=10:5 wt%] (c) SEM images of GH₁₅₀. [The preparation of GH₁₅₀ was consistent with GH₁₀₀ whereas the concentrations of GO-HAP were 150 ng mL⁻¹.] (d) SEM images of GH₂₀₀. [The preparation of GH₂₀₀ was consistent with GH₁₀₀ whereas the concentrations of GO-HAP were 200 ng mL⁻¹.]

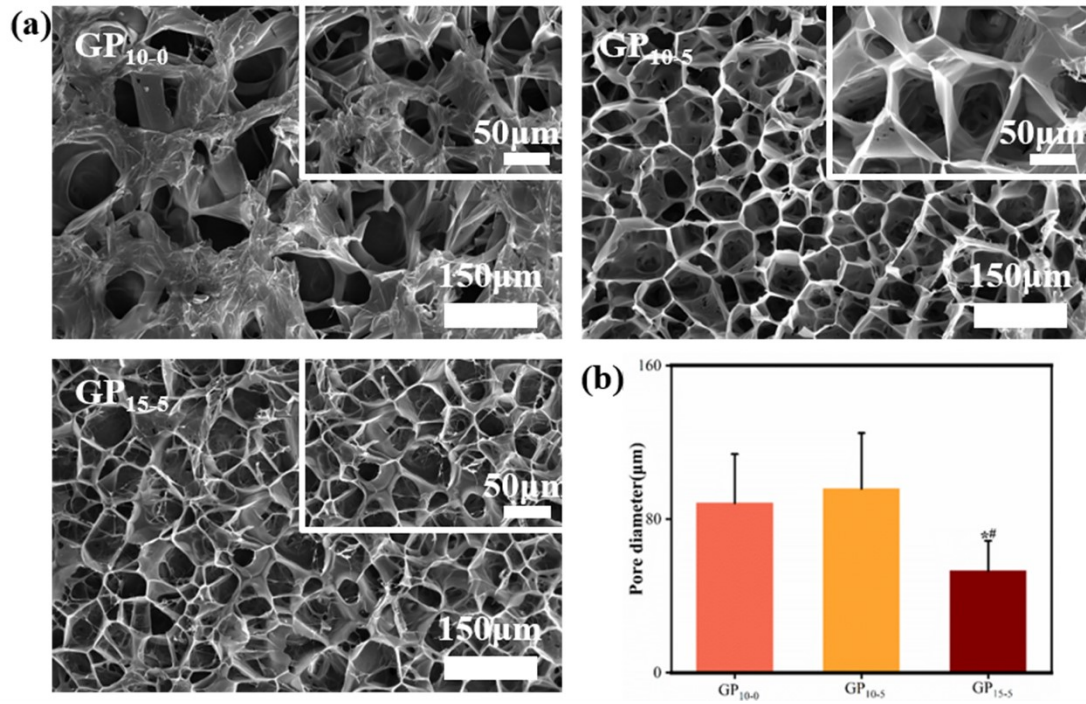


Fig. S5. (a) SEM images of: GP₁₀₋₀, GP₁₀₋₅ and GP₁₅₋₅ at high and low magnifications. Scale bars = 50 μm and 150 μm. (b) The pore diameters of GP₁₀₋₀, GP₁₀₋₅ and GP₁₅₋₅, * and # represented $p < 0.05$ when compared with GP₁₀₋₀ and GP₁₀₋₅, respectively.

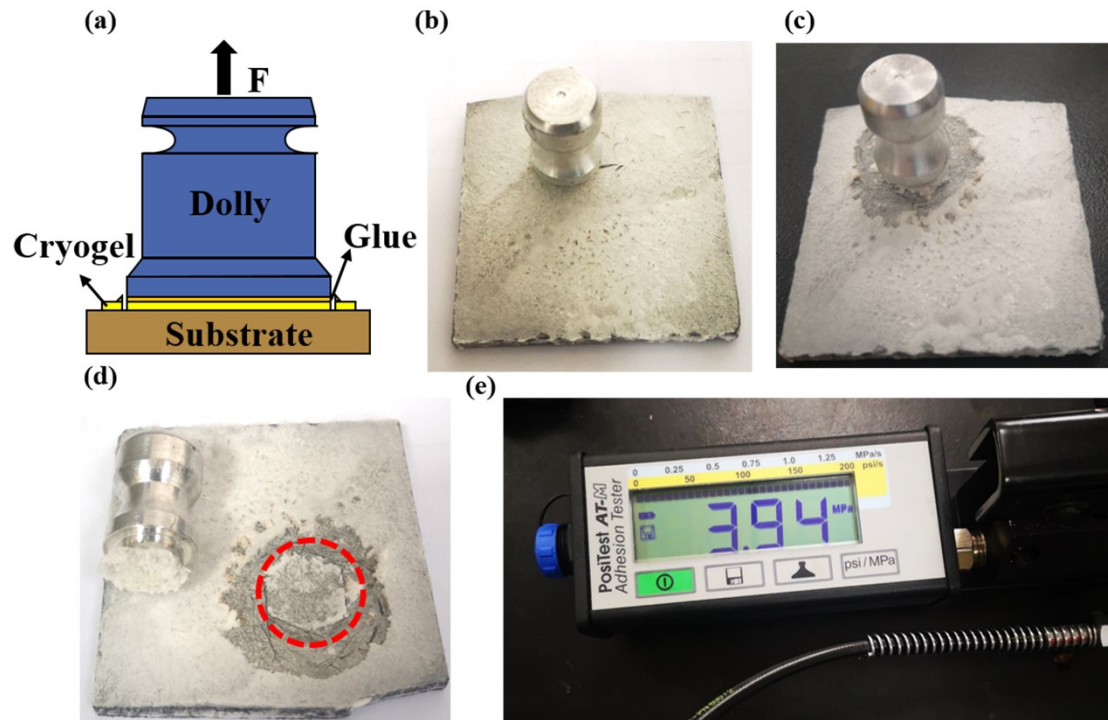


Fig. S6. (a) Diagram of the experimental device of the pull-off method. (b) Image of GP₁₀₋₅ after adhering the dolly with epoxy glue. (c) Image of GP₁₀₋₅ after cutting by the cutter. (d) Image of GP₁₀₋₅ and the dolly after pull-off test. (e) Adhesion data of GP₁₀₋₅.

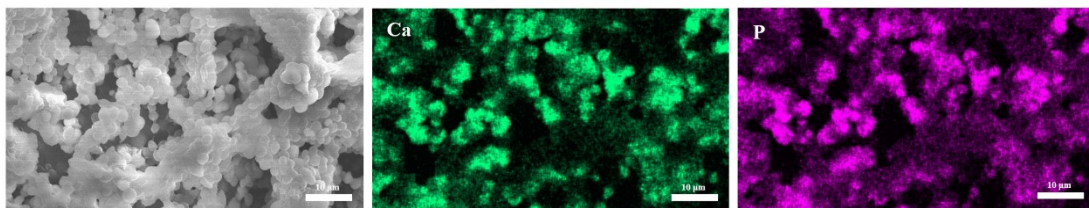


Fig. S7. EDS images of GH₁₀₀ after soaking in SBF for 7 days. Scale bars = 10 µm.

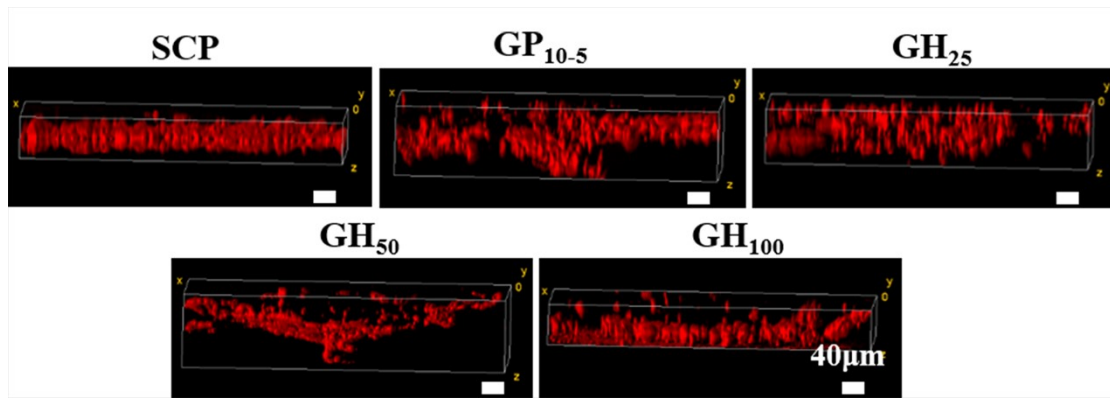


Fig. S8. Representative 3D CLSM images of HUVEC migration into different scaffolds stained using TRITC-phalloidin for cytoskeletal organization. Scale bars = 40 μm .

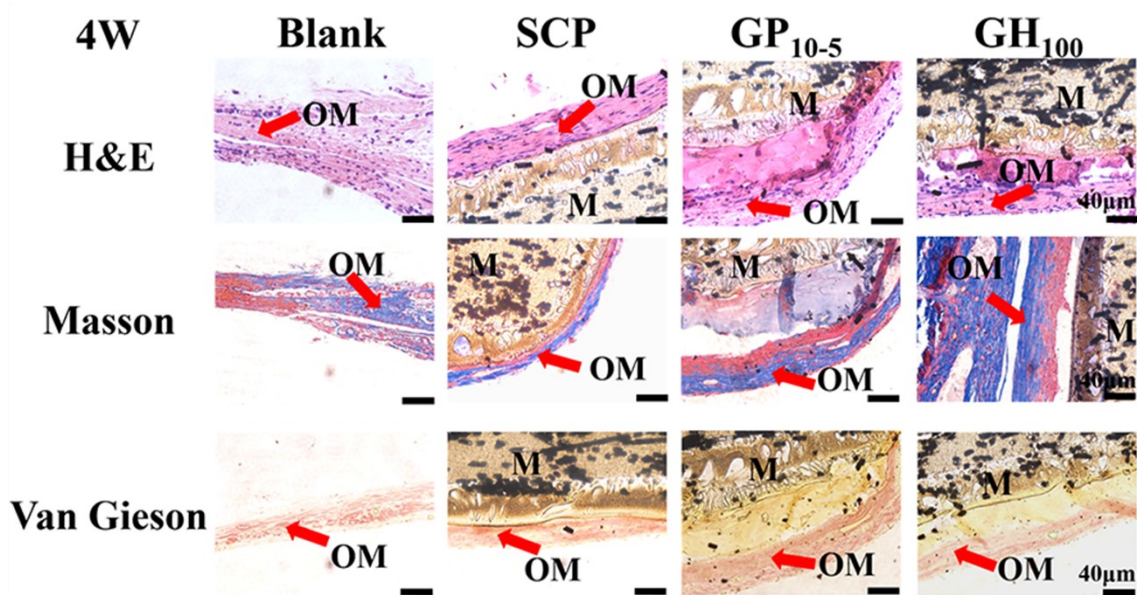


Fig. S9. H & E, Masson and Van Gieson staining of the calvarial defect regions after treatment with different samples for 4 weeks. Scale bars = 40 μm . [M, Materials; OM, osteoid matrix (red arrows)].

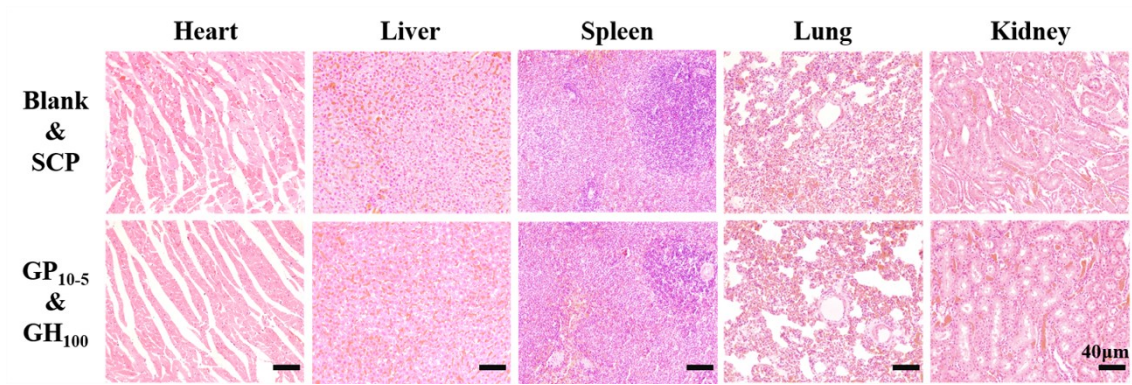


Fig. S10. Histological observation of H & E staining of tissues (heart, liver, spleen, lung, and kidney) after 12 weeks of treatments. Scale bar = 40 μ m.

Table S1 Primer pairs used in RT-PCR analysis.

Genes	Forward primer sequence (5'-3')	Reverse primer sequence (5'-3')
BMP-2	ATTAGCAGGTCTTTGCACCA	ACGCTTTTCTCGTTTGTGGA
Runx2	GGCAGCACGCTATTAAATCCA	GACTCATCCATTCTGCCGCTA
	A	
ALP	ATCGGAACAACCTGACTGACC	CTGCCTCCTTCCACTAGCAA
COL-I	ATGCCATCAAGGTCTACTGCAA	GAACCTTCGCTTCCATACTCG
GAPDH	TATGACTCTACCCACGGCAAG	ATACTCAGCACCAGCATCACC

Table S2 Elemental composition of different substrate surfaces as determined by XPS.

Material	C (%)	N (%)	O (%)	S (%)	Ca (%)	P (%)
SCP	58.84	---	40.06	1.09	---	---
GP ₁₀₋₅	41.17	10.11	48.72	---	---	---
GH ₁₀₀	44	12.46	40.97	---	1.6	0.96

Table S3 Changes in Ca and P content of GH₁₀₀ sample before and after immersion in SBF by EDS.

Sample	Ca (%)	P (%)	Ca/P
Before immersion in SBF	1.6	0.96	1.67
After immersion in SBF	3.87	2.1	1.84
New bone-like apatite	2.27	1.14	1.99