

Electronic Supplementary Information

Characterization of organophosphatic brachiopod shells: spectroscopic assessment of collagen matrix and biomineral components†

Oluwatoosin B. A. Agbaje,^{*a-c} Simon C. George,^b Zhifei Zhang,^d Glenn A. Brock,^{c,d} Lars E. Holmer^{a,d}

^a Department of Earth Sciences, Palaeobiology, Uppsala University, Uppsala, Uppsala, Sweden

^b Department of Earth and Environmental Sciences and MQ Marine Research Centre, Macquarie University, Sydney, Australia.

^c Department of Biological Sciences, Macquarie University, Sydney, Australia.

^d State Key Laboratory of Continental Dynamics, Shaanxi Key Laboratory of Early Life & Environments, Department of Geology, Northwest University, Xi'an, 710069, China.

*** Correspondence:** toosin.agbaje@mq.edu.au; toosin91014@gmail.com

SI Table 1a. Wavenumber from Raman and FTIR spectra of type I collagen and hydrogen peroxide-treated brachiopod shells.

Type 1 Collagen		Chondroitin sulfate A		<i>D. tenuis</i>		<i>L. anatina</i> (ML/RL)		Assignment/Component
Raman	FTIR	Raman	FTIR	Raman	FTIR	Raman	FTIR	
1668				1664				Amide I, β -sheet/ β_{10} helix, Collagen
1657						1654		Amide I, α -helix, GAGs/Collagen
	1634		1635*		1637		1634/6	Amide I, Triple helix, Collagen/GAGs
*1616				1616		*1616		Amide I, Collagen
1604			1606	1606		1604		Amide I
1584				1585		1585*		ν C=C ring of phenylalanine
		1572						GAGs
1555	1544		1566*	1557	1538	1543*	1538/44	Amide II
1451	1450	1448*		1451	1455	1452	1454/47	ω CH ₂ / δ asCH ₃ /Phospholipids
1423					1417	1422		ν COO ⁻
1398	1401	1414	1411				1403/1395	
		1375	1375	1384*	1380	1379w		ν_s CH ₃ of GAGs
1336/43	1337	1341		1337	1338	1336	1337/8	Amide III, α -helix, Collagen/GAGs
1317	1317	1316*	1310	1319	1318	1313	1317/5	Amide III, α -helix
	1281	1277		1289w	1282	1286	1282	Amide III, α -helix
1265		1267*		1265		1264		Amide III, α -helix, Collagen/GAGs
1245	1235	1237*		1242	1237	1238/42	1236	Amide III, β -sheet and/or random coils,
			1226/55					Sulfate asymmetric stretching of GAGS
1208		1201w		1205		1206/7		ν C-C of tryptophan and phenylalanine
	1202				1202		1203	wCH ₂ from Collagen
1165/74				1172		1174		ν C-O-C, Collagen
	1161	1151*	1155	1157	1156	1155	1158/60	ν C-O-C of polysaccharide (GAGs)
1125		1138	1123	1126		1134	1125/6	ν C-C of collagen/C-O-S of GAGs overlaps with the HPO ₄ ²⁻
1097		1090		1104	1108*	1107	1116	ν_1 CO ₃ ²⁻ ; A-type
1086*	1080			1084		1086*	1080	ν C-O in collagen and GAGs overlaps with ν_3 PO ₄ ³⁻
1078*				1077*		1072/4		ν_1 CO ₃ ²⁻ ; B-type
1067	1061	1069	1062*	1064*	1065*		1069*/4	ν C-C-O of carbohydrate in collagen and GAGs overlaps with lipids
		1055				1053		GAGs overlap with ν_3 PO ₄ ³⁻
1046				1040		1040		ν C-O of carbohydrate in collagen and GAGs/ ν_3 PO ₄ ³⁻
1031	1031		1027	1032	1025	1032	1035/6	ν C-O of protein; ν_3 PO ₄ ³⁻ overlaps with proline ν C-C
		1019		1012*		1011*		GAGs overlap with HPO ₄ ²⁻
1003				1004		1002/3		ν C-C of phenylalanine (Collagen)
965	971	978	985*	964	952	964/5	968/71	ν_{as} PO ₄ ³⁻ (hydroxyapatite); ν C-O-S
				942	944*	942	943/4	ν C-C backbone of collagen
939	937	939	937*	939		936		ν C-O-C backbone of collagen/GAGs
920	922		924	921				ν C-O-C backbone of collagen/GAGs
						905		ν C ^a -C, ν C-N, ν CH ₃
		885	891*	887			895/7	δ C-H for anomers (GAGs)
874	875			877	870	872/3	875/3	ν C-C of hydroxyproline
853	852	855	855	855	856*	852/3		ν C-C of proline (Collagen)/GAGs
				827/33				ν C-COO ⁻ of tyrosine (Collagen)
815	815			818	810*	814	814/9	ν C-O-C collagen crosslink
785		783	793*	778		786/76		ν (C-C)/pyrimidine ring breathing
759		760		758		758/60		δ C-COO ⁻ / ν_4 CO ₃ ²⁻ ; B-type
		722	726	714, 701		730, 704		ν_1 CO ₃ ²⁻ ; B-type ν_1 CO ₃ ²⁻ ; B-type
		693		671		679/7		ν_4 CO ₃ ²⁻
		642	655	642		643		τ C-C, ν COO ⁻ of Collagen/GAGs

621				621		619		<i>t</i> C-C Collagen
				602		607		$\nu_4\text{PO}_4^{3-}$
590*		590		591 579		591, 580		$\nu_4\text{PO}_4^{3-}$ overlaps with GAGs
563		547				556		$\nu(\text{S-S})$ in collagen/GAGs
534				536		527		$\nu(\text{S-S})$ in collagen
476		461/92		476		470		$\nu(\text{S-S})$ in collagen/GAGs
				450		450/4		$\nu_2\text{PO}_4^{3-}$
		439		431		431		$\nu_2\text{PO}_4^{3-}$ overlaps with GAGs
		410						GAGs

Notes: modern/living (ML) and recent (RL) *Lingula anatina* shells. Chondroitin sulfate A represents sulphated glycosaminoglycan

SI Table 1b. Raman and FTIR band assignment for pure α -chitin and chitin (organic) fibre

α -chitin		Chitosan		Assignment
Raman	FTIR	Raman	FTIR	
1655	1656	1657	1651	Amide I, α -helix
1617	1621	1598	1590	Amide I, ν C=O
	1553		1562	Amide II
1447		1460		CH ₂ , CH ₃
	1428		1422	CH ₂
1413	1415	1410		ν sCOO ⁻
1376	1376	1374	1376	r C-CH ₂ , δ C-CH ₃
1327		1323	1315	Amide III, ν CH ₃
	1308			Amide III, CH ₂ w,
1265	1260	1263	1260	Amide III, ν C-H, δ N-H, δ C=O
	1234	1225	1235	Amide III, N-H ib, C-N s
1202	1204	1202	1202	Amide III, C-O-H d
1147	1154	1146	1151	ν C-O-C
1111/04	1113	1111		ν C-O-C, ring
	1068	1087		ν C-O-H
1057			1061	ν C-O, ν C-C, ring
1031	1024	1036	1027	δ C-O-H, ν C-O, ν C-C skeletal
	1010	989	996	δ C-O-H, ν C-O
971*				δ CH ₃ , ν C-O
953	952	942	951	δ CH ₃ , δ C-O-H
915	919			CH ₃
895	895	895	895	δ CH _x , C-O-C glycosidic bond
				C-C, C-O, C-H
				C-C, C-O, C-H
	746			CH ₂
710	701	707		C-O, C-H, N-H
	692			C-O, C-H, N-H
648	634		660/7	C-C, C-O, C-H
598				C-C, C-O, C-H,
				PO ₄ ³⁻ s
563		570		C-C, O-H
530				C-C skeletal backbone
496		495/78		C-C skeletal backbone
450/5		441		C-C-C ring
		420		C-C-C ring

SI Table 2. Assignment of amide I peak positions and percentage areas to secondary structure.

Sample			
	FTIR	Area %	Assignment
Type 1 Collagen (TC)			β -turn
	1692	4	β -turn
	1661	51	β -sheet, 3_{10} helix
	1651	12	α -helix
	1644	4	Unordered
	1636	6	3_{10} -helix
	1629	23	3_{10} -helix
	1616		Side chain, Tyrosine and tryptophan
Recent <i>D. tenuis</i> (DT)			β -turn
	1692	3	β -turn
	1678	13	β -sheet
			β -sheet
	1660	34	β -sheet, 3_{10} -helix
	1650/8	2	α -helix
	1643	22	Unordered
	1631	26	3_{10} -helix
			β -sheet
			β -sheet; Side chain
	1616		Side chain, Tyrosine and tryptophan
	1603		Side chain, tyrosine
	1595		Side chain, Phenylalanine
Modern <i>L. anatina</i> (ML)			
			β -turn
	1692	2	β -turn
	1680	7	
	1662	23	β -sheet, 3_{10} -helix
	1652	29	α -helix
			Unordered
	1635	32	3_{10} -helix
	1626	7	β -sheet
	1616		Side chain, Tyrosine and tryptophan
	1603		Side chain, tyrosinate

	1591		Side chain, Phenylalanine
Recent <i>L. anatina</i> (RL)			β -turn
	1689	6	β -turn
			β -turn
	1660	54	β -sheet, 3_{10} -helix
	1652	6	α -helix
	1640/ 6	5	Unordered
	1631	29	3_{10} -helix
			β -sheet
	1615		Side chain, Tyrosine and tryptophan
	1600		Side chain, tyrosinate
	1591		Side chain, Phenylalanine

Notes: ML = modern, RL = recent.

SI Table 3. A. Raman spectra (cm^{-1}) peak position of the phosphate group in brachiopod shells.

<i>D. tenuis</i>		<i>L. anatina</i> (ML)		<i>L. anatina</i> (RL)		Component
Shift	FWHM	Shift	FWHM	Shift	FWHM	
948	7	949	8	950	14	ACP [1]
955	8	955	7	955	8	OCP [1, 2]
958	2			959	3	CAP [2, 3]
964	11	964	10	965	11	HAP [2, 3]
973	7	971	12	975	10	TCP [1, 3]
~980	5					DCPD [3]

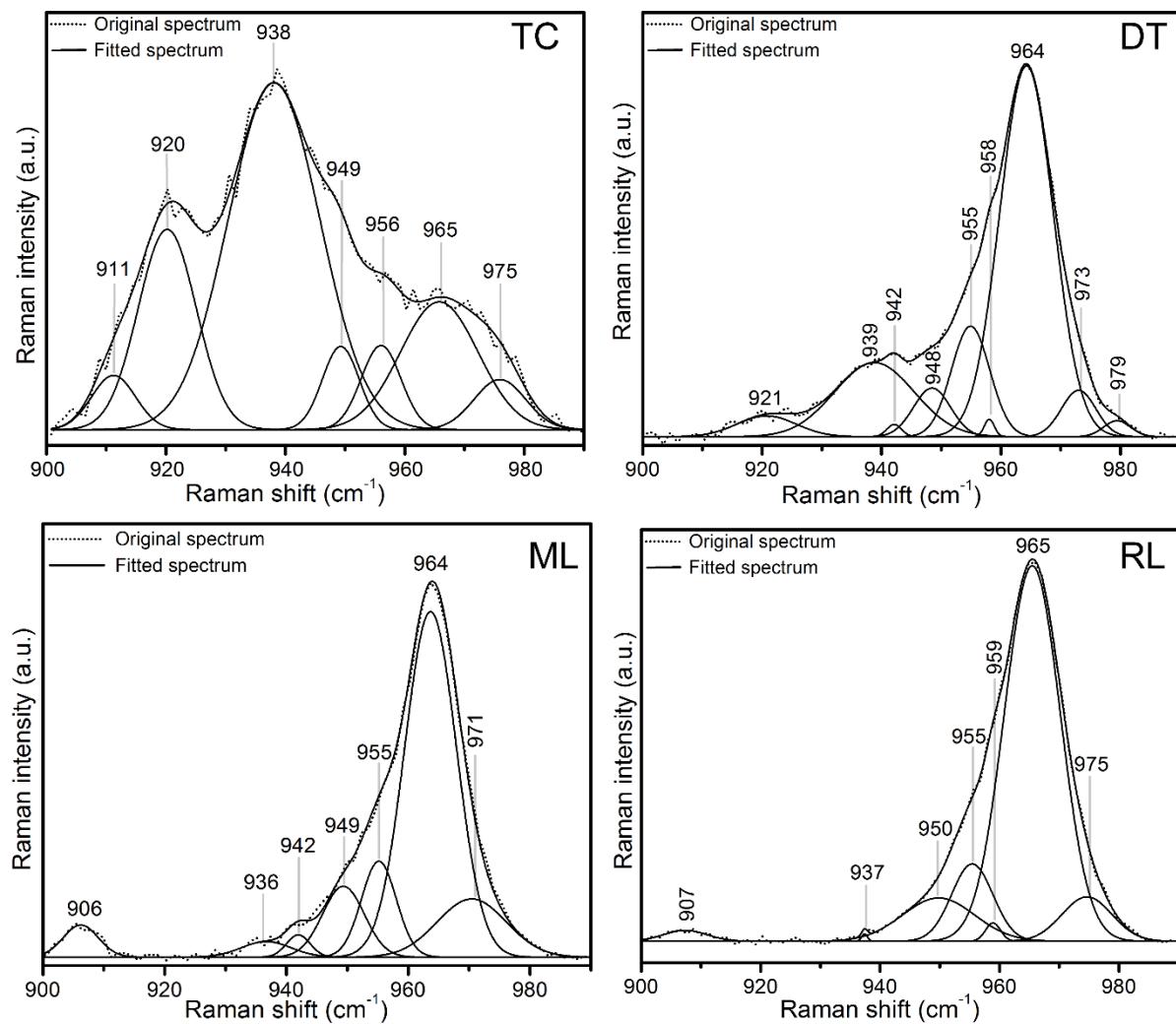
Notes: ML = modern, RL = recent, FWHM = full width measured at half maximum intensity, ACP = amorphous calcium phosphate, OCP = octacalcium phosphate, CAP = carbonated hydroxyapatite, HAP = hydroxyapatite, TCP = tricalcium phosphate, DCPD = Dicalcium phosphate dihydrate.

SI Table 4. Collagen crosslinks 1660/1690 ratio of type I collagen and brachiopod shells.

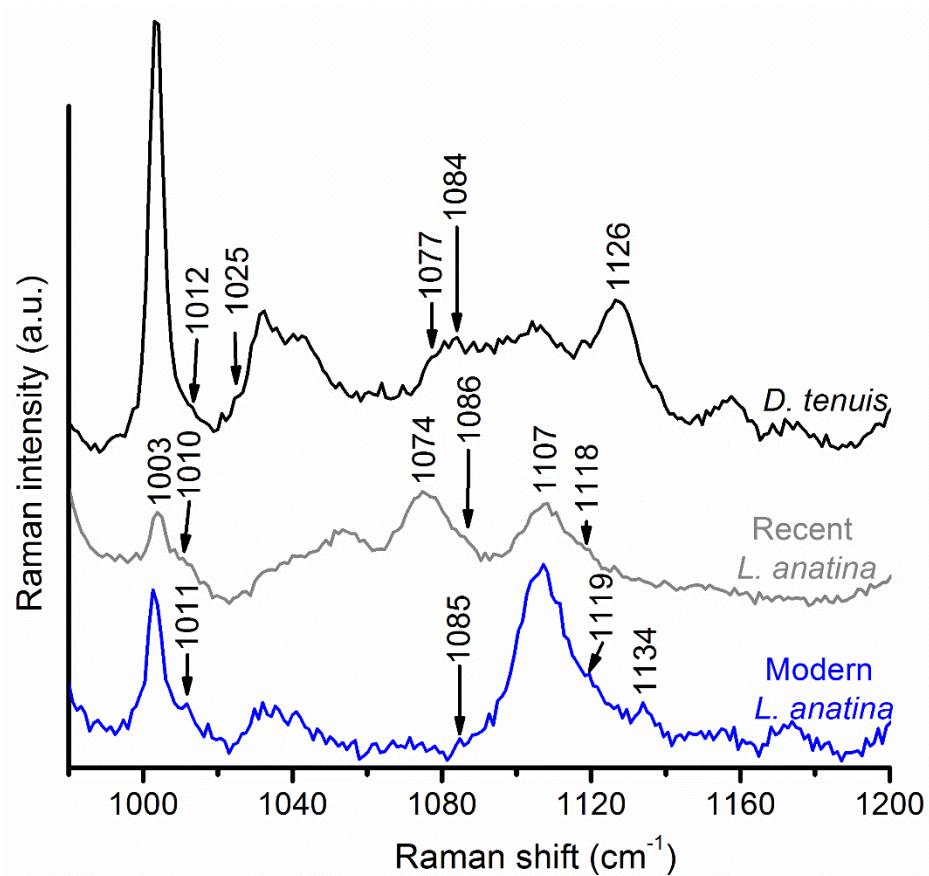
	Type I collagen	<i>D. tenuis</i>	<i>L. anatina</i> (ML)	<i>L. anatina</i> (RL)
Collagen cross-links	13.0±0.53	11.5±0.47	11.0±0.42	9.9±0.39



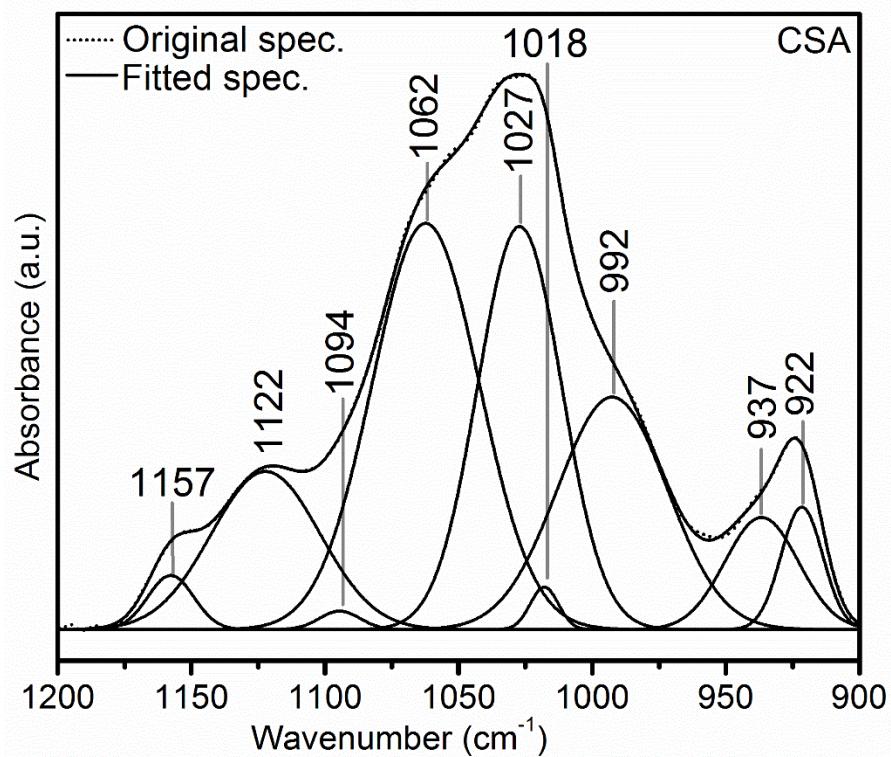
SI Figure 1. *Lingula anatina* and *Discinisca tenuis* showing the interior part of the shells. The white dashed line in *Lingula anatina* and the black dashed region in *Discinisca tenuis* indicate part of the shell that was exempted from vibrational spectroscopy analyses and SEM imaging. Black dashed line in *Lingula anatina* denotes the region that was measured for microRaman and Fourier Transform Infrared.



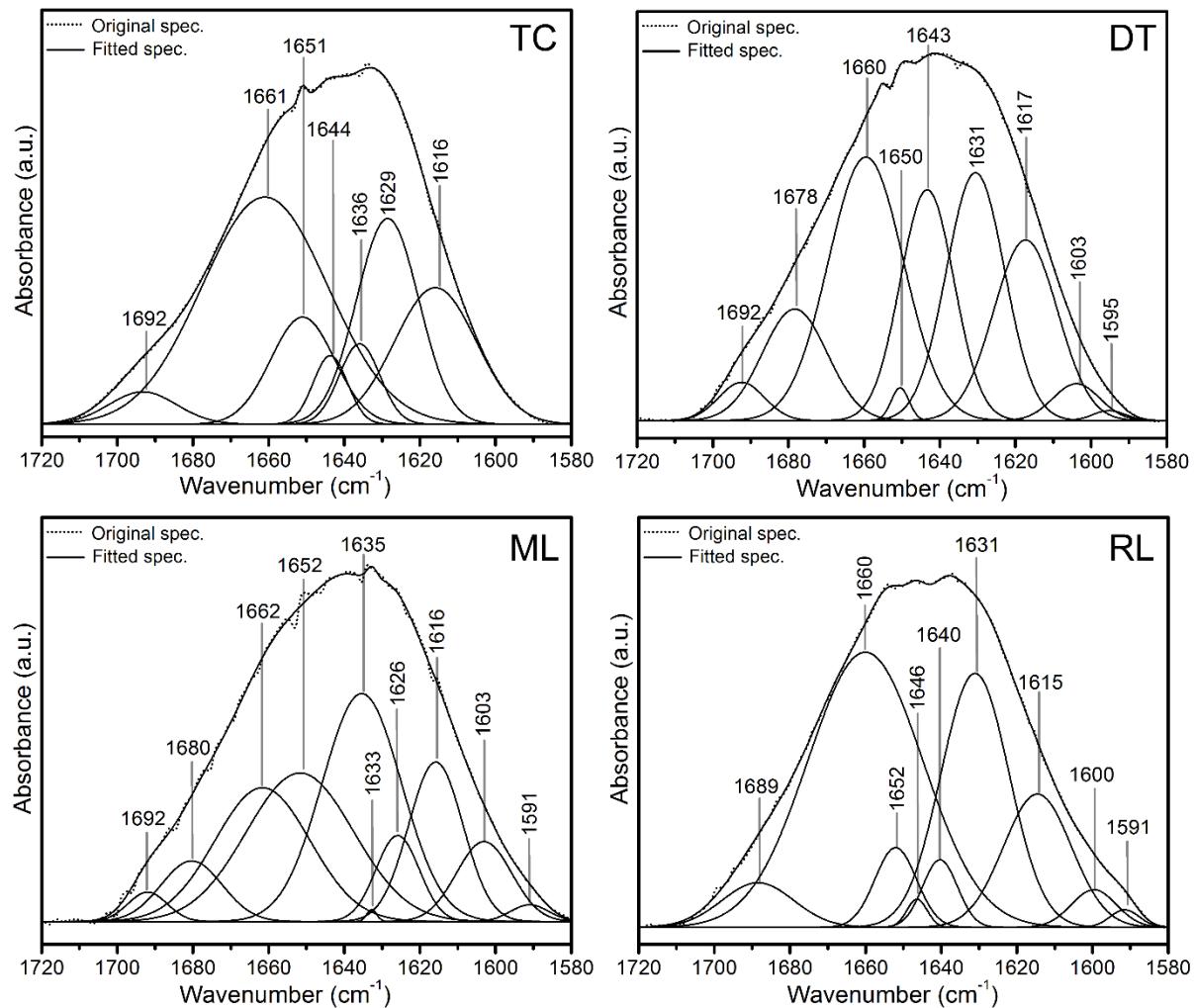
SI Figure 2: Raman spectra of type I collagen (TC) and shells of brachiopods (DT, ML and RL) in the 900–990 cm^{-1} region. The Raman data were deconvoluted by a Gaussian fit. See text for the interpretation of the peaks. DT, ML and RL represent *Discinisca tenuis*, recent *Lingula anatina* and modern/living *Lingula anatina*, respectively.



SI Figure 3: Raman spectra of brachiopods shells in the 980–1200 cm⁻¹ region.



SI Figure 4. Curve-fitting of FTIR spectrum of Chondroitin sulphate A (CSA; glycosaminoglycan) in the 1200–900 cm⁻¹ region. See **Table 2** for peak assignments.



SI Figure 5: Curve-fitting of FTIR spectra of amide I in the 1720–1580 cm⁻¹ region. TC, DT, ML and RL represent type I collagen, *Discinisca tenuis*, recent *Lingula anatina* and modern/living *Lingula anatina*, respectively.

References

- 1 N. J. Crane, V. Popescu, M. D. Morris, P. Steenhuis, M. A. Ignelzi Jr, *Bone*, 2006, **39**(3) 434-442.
- 2 J. Freeman, B. Wopenka, M. Silva, J. Pasteris, *Calcif. Tissue Int.* 2001, **1**, 68(3).
- 3 S. Koutsopoulos, *J. Biomed. Mater. Res. A*, 2002, **62** 600-612.