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Mesoporous Materials Modified by Aptamers and Hydrophobic Groups Assist Ultra-sensitive Insulin Detection in Serum

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ABSTRACT: A novel mesoporous material modified with both insulin-binding-aptamer and hydrophobic methyl groups is synthesized. With rational designed pore structures and surface chemistry, this material is applied in sample pre-treatment for ELISA, enables a quantification (0.25-5 pg ml⁻¹) of insulin in serum, 30-folds enhancement of limit-of-detection comparing to commercial ELISA kit.

Detection of insulin is essential for disease diagnosis,¹ pathological research and doping control.^{2, 3} Due to low abundance and complexity in biological samples (e.g. serum),⁴ sensitive detection of insulin remains a long-standing challenge. Many strategies such as enzyme linked immunosorbent assay (ELISA), radioimmunoassay (RIA)³, immunoassay followed by liquid chromatography–tandem mass spectrometry (LC-MS/MS)⁵, solid phase extraction (SPE),⁶, ⁷ photolytic-electrochemical detection⁸ and high performance liquid chromatography(HPLC) coupled with UV absorbance^{9, 10} have been developed for insulin detection. However, for detection in complicated samples containing very low levels of insulin,^{3, 11} it is essential to pre-treat and enrich samples to increase the detection sensitivity.^{12, 13}

Recently, the application of nanotechnology has led to significant advances in biomolecule enrichment and identification.^{14, 15} In particular, mesoporous materials with controllable pore structure and surface chemistry have attracted much attention.^{13, 14} Our previous study showed that hydrophobic modification of pore surface with a suitable pore size is beneficial to insulin enrichment.¹³ However, the advantages provided by optimised pore size and hydrophobicity do not address the specific binding issue. It is expected that mesoporous materials functionalized with specific binding molecules will minimise non-specific adsorption in serum and selectively enrich low abundance targets.

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Antibody is a well-known specific binding molecule with high affinity of antigen-antibody binding. There are several reports on mesoporous silica materials functionalized with antibodies (generally on the external surface) for protein captur. Despite the stability issue,¹⁷ it is difficult to modify antibod inside the mesopores due to the large molecular size (e.g. IgC 20 nm).¹⁸ Although large-pore materials are available,¹⁹ previous study has shown that a large pore size is not favoure 1 for the insulin enrichment.¹³ Alternatively, aptamers are effective substitutes for antibodies¹⁷ and have advantages such as small molecular size, excellent stability, and low cost.²⁰ It is reported that the insulin binding aptamer (IBA) with a tworepeat parallel G-quadruplex structure can selectively bind to insulin.^{20, 21} A study used IBA modified plate for insulin detection, achieving a limit-of-detection (LOD) in standard solution and serum of 5 and 20 ng ml⁻¹, respectively.²² IBA w(5 modified onto carbon nanotubes for insulin extraction from pancreatic INS-1 cells with an LOD of 58 ng ml-1.20 However there is no report on the modification of mesoporum materials by IBA for insulin detection. Furthermore, although aptamers have been modified onto mesoporous materials for drug delivery,²³ there is no report on co-modification of hydrophobic groups and aptamers for insulin enrichment.

In this study, the conjugation of IBA and methyl groups on mesoporous silica is reported for sensitive insulin detectio . Aminopropyl(diethoxy)methylsilane (APDEMS) is used t modify mesoporous silica to introduce two functional groups , CH₃ and -NH₂ for further IBA conjugation) simultaneously o one silanol (Scheme 1a), which overcomes the challenge typically encountered in two-step modification processes whereby the first modification step consumes a large number of available silanols and reduces the density of the second functionality.²⁴ The resultant functional mesoporous silic enables efficient enrichment of insulin in serum (Scheme.1r prior to ELISA test, achieving a quantification range of 0.25pg ml⁻¹ of insulin in serum, 30 folds enhancement compared tr commercial ELISA kits alone and more sensitive than othe literature reports (Table. S1).⁵⁻⁷, 9, 10, 20, 22, 25-30

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Scheme 1 Schematic illustration showing the (a) synthesis process of insulin binding aptamer and hydrophobic modified mesoporous materials for insulin enrichment; (b) process of pre-treatment coupling ELISA. * HRP: Horseradish peroxidase.

To demonstrate the effect of surface chemistry on insulin enrichment, three functionalized mesoporous silica materials were prepared using SBA-15^{31, 32} as the parent material. The first one contained both -CH3 and -NH2 groups (named as SBA-15-CH₃-NH₂). The second material was prepared with -NH₂ but without -CH₃ group (SBA-15-NH₂) in order to compare the effect of hydrophobic modification. The third one was prepared with both -CH₃ and -NH₂ groups on the outer surface [SBA-15-CH₃-NH₂ (out)] in order to confirm the surface modification inside the mesopores. After linking IBA to -NH₂, the corresponding materials are named as SBA-15-CH₃-IBA, SBA-15-IBA and SBA-15-CH₃-IBA (out).

The small-angle X-ray scattering (SAXS) patterns of SBA-15 (Fig. S1a), SBA-15-NH₂ (Fig. S1b), SBA-15-CH₃-NH₂ (Fig. S1c) and SBA-15-CH₃-NH₂ (out) (Fig. S1d) are shown in Fig. S1. In each pattern, three well-resolved diffractions can be indexed to the 100, 110, and 200 reflections of an ordered two-dimensional hexagonal mesostructure. For SBA-15, SBA-15-CH₃-NH₂ and SBA-15-NH₂, the diffractions appear at the same position corresponding to a cell parameter (a) of 11.3 nm, suggesting that the nanostructure is well maintained after modification. For SBA-15-CH₃-NH₂ (out), the 100 diffraction shifts to the small angle range with an increased *a* of 11.9 nm. Moreover, the 200 diffraction has an obviously higher intensity compared to 110 peak (Fig. S1d). The above observation is attributed to the difference in sample preparation. Compared to the first three samples based on calcined SBA-15, SBA-15-CH₃-NH₂ (out) was prepared starting from as-synthesized SBA-15 followed by modification and surfactant extraction. The silica matrix shrinkage is less during the extraction process, leading to a slightly larger unit size. The change in relative intensity between 110 and 200 diffractions can be attributed to the the thickness of silica walls as reported previously.³³

Transmission electron microscopy (TEM) images in Fig. S2 display the ordered pore channels of SBA-15 (Fig. S2a) with different modifications, including SBA-15-CH₃-NH₂ (Fig. S2b), SBA-15-NH₂ (Fig. S2c) and SBA-15-CH₃-NH₂ (out) (Fig. S2d). Nitrogen adsorption isotherms for all silica materials an typical type IV with capillary condensation of nitroge occurring at a relative pressure (P/P0) in the range of 0.60 an 0.80. The Barrett–Joyner–Halenda (BJH) model was used t calculate the pore size distribution curves from the adsorptic branch (Fig. S3, Table 1). The pore sizes of SBA-15, SBA-15-CH₃-NH₂, SBA-15-NH₂ and SBA-15-CH₃-NH₂ (out) were measured to be 8.0, 6.68, 6.70, and 7.1 nm (Table 1), respectively. Decreases in pore size are observed for modified materia s when compared to pristine SBA-15 (pore size 8.0 nm) due to the occupation of the pore by modified groups. The fourth material (SBA-15-CH₃-NH₂ (out)) also showed a decrease ... pore size compared to the pristine SBA-15 (8.0 to 7.1 nm), bu . not as much as SBA-15-CH₃-NH₂ (6.68 nm) and SBA-15-NH (6.70 nm), consistent with SAXS results. The surface area an pore volume of all the materials are listed in Table 1.

FTIR spectroscopy was used to confirm modifications (S4), revealing two peaks at around 2900 cm⁻¹ which indicate the successful incorporation of -CH₃ groups (Fig. S4). potential measurements show that the pristine SBA-15 (Table. 1) is negatively charged due to deprotonation of the group. For all the modified materials, the positive ζ potentia suggest the successful modification with -NH₂ group.

For the conjugation of IBA and nanomaterials, we followed well-developed method which used the N-(?-Dimethylaminopropyl)-N'-ethylcarbodiimide hydrochlorica (EDC) as a reagent.^{35, 36} In order to characterize materials after IBA functionalization, TEM images for SBA-15-CH₃-IBA weil recorded (Fig. 1), which show ordered pore channels from the 100 (Fig. 1b) and 110 (Fig. 1b) directions, confirming that the originally ordered SBA-15 pore structure remains aft aptamer modification. We did not conduct the nitroge. adsorption test (each test usually requires a minimum of 5 mg sample) for IBA modified materials due to the relative hig. price of IBA (even it is cheaper than antibodies).

Table 1. Pore size, surface area, pore volume and zeta potential of materials

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Sample	Pore size	SBETa	Vpb	ζ	
	(nm)	(m2/g)	(cm3/g)	potential	
SBA-15	8.00	578	0.98	-17.5	
SBA-15-CH3-NH2	6.68	297	0.53	35.6	
SBA-15-NH2	6.70	286	0.46	36.7	
SBA-15-CH3-NH2(out)	7.10	302	0.51	30.6	

^aBET surface area. ^bTotal pore volume.



Fig. 1 TEM images of SBA-15-CH₃-IBA viewed (a) parallel and (b) perpendicular to the channel direction of SBA-15. Scale bar: 50 nm.

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The surface modification of various groups was also confirmed by X-ray photoelectron spectroscopy (XPS) analysis. Compared to unmodified SBA-15 (Fig. 2a), the existence of nitrogen and an increased amount of carbon in the case of SBA-15-CH₃-NH₂ (Fig. 2b) suggest the successful functionalization of both -CH₃ and -NH₂. XPS survey was also conducted to investigate the modification of IBA. The carbon spectrum of SBA-15-CH₃-NH₂ (Fig. 2c) can be well-fitted to two peaks at 284.8 and 286.1 eV, which are attributed to the binding energies of carbon atoms bonded to C/Si and N, respectively. For SBA-15-CH₃-IBA (Fig. 2d), the spectrum reveals four peaks at 284.8, 285.7, 286.8 and 288.6 eV, which correspond to the binding energies of C-C/C-Si, C-N, C=O and N-C=O respectively. The existence of C=O suggest the successfully modification of IBA.

To find out the optimised material for insulin enrichment, the enrichment efficiency of all groups were compared. The biological system selected for testing was horse serum. In order to remove interferences arising from non-target components in the complex bio-sample, a prior purification step using MOSF-CH₃ was applied using a literature protocol to remove large proteins before the enrichment step.¹³ Fig. S5a displays the MS spectrum obtained from the untreated insulin solution (10 ng ml⁻¹). No signal can be detected due to the low insulin concentration. When applying the designed SBA-15-CH₃-IBA for insulin enrichment in a low concentration of 0.05 ng ml⁻¹, both single and double charged peaks are observable (Fig. S5b). For the pre-treatment with SBA-15-IBA and SBA-15-CH₃-IBA (out), the detection limits are 1 ng ml⁻¹ (Fig. S5c) and 0.5 ng ml⁻¹ (Fig. S5d), respectively. The results show that the SBA-15-CH₃-IBA material (Fig. S5b) has the highest detection sensitivity among all the groups due to the specific binding from IBA, hydrophobic interaction from -CH₃ and size selective effect from mesopores.



Fig. 2 XPS survey scan of a) SBA-15 and b) SBA-15-CH3-NH2; the corresponding fine spectra of c) SBA-15-CH3-NH2 and d) SBA-15-CH3-IBA.

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The initial results indicate that SBA-15-CH₃-IBA with both aptamer and hydrophobicity modification is the best material for insulin enrichment. The design SBA-15-CH₃-IBA (Fig. S51) has better performance comparing to SBA-15-IBA (Fig. S⁻c) which suggests that the hydrophobicity plays an important role in the insulin enrichment. The conclusion is consistent with previous reports.^{13, 37} The modifications of IBA and -CH₃ into mesopores enhanced the enrichment efficiency (0.05 ng ml⁻¹) since the outer modified group has a worse detection limit (0.5 ng ml⁻¹). The reason is more IBA and CH₃ can be modified into nanoparticle when the inner surface is accessible. The detection limit of 0.05 ng ml⁻¹ in serum is mucnlower than in most of the non-antibody based (116-290 ng m ¹)³⁸⁻⁴⁰ or aptamer based (20-58 ng ml⁻¹)^{20, 22} enrichments, an comparable to antibody based detection.³⁰

After confirming the designed SBA-15-CH₃-IBA is the optim. material for insulin enrichment, a further application u enrichment to enhance ELISA was conducted. Fig. 3a shows the standard curve of a commercial ELISA kit, which obtained by testing the standards in this kit following the instruction provided by manufacturer (Scheme S1). All standards in this kit are in human serum, and the quantification limit is 7.7 pg ml⁻¹. To achieve a better detection efficiency and conduct the quantification analysis in lower concentration, the designed SBA-15-CH₃-IBA was applied to pre-treat samples. A series of diluted insulin standard solution. with ultra-low concentration (0.25-5 pg ml⁻¹) were tested t ELISA after pre-treatment. The quantification in lo... concentration range (0.25-5 pg ml⁻¹) is achieved as shown i the standard curve (Fig. 3b) of enhanced ELISA. The linea. regression coefficient (R²) is 0.9556, which is not perfect line but sufficient for quantification in low concentration range. (pg ml⁻¹). The results indicate that with the help of designe SBA-15-CH₃-IBA, the detection sensitivity of commercial ELIS, kit increased. The limit-of-detection (0.25 pg ml⁻¹) is 30 times better than that of commercial ELISA kit (7.7 pg ml⁻¹).

To further confirm the quantification ability, a standard addition technique was employed and a recovery rate was calculated (see SI).³⁵ In our case, a recovery rate of 71% with achieved, which is comparable to the recovery rates is previous studies generally conducted at high concentrations.²⁷ ³⁸ For example, a study used self-assembled TiO₂ nanocrystriculaters to enrich beta-casein (0.75mg ml⁻¹) and achieved recovery efficiency of 50%.³⁶ In another study, the recover rate of glycopeptide (1µg µl⁻¹) enriched by functionalized magnetic nanoparticles is 77.8%.³⁷ By comparing with literatures, a recovery rate of 71% in the very low concentration of 0.25-5 pg ml⁻¹ is significant.



Fig. 3 Standard curves showing the quantification range of (a) commercial ELISA kit, (t ultrasensitive ELISA with pre-treatment by combo-pore-CH₃-IBA.

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In conclusion, we have successfully synthesized mesoporous nanomaterials with co-modified aptamer and methyl groups. When applying this material in the combo-pore approach⁷ for pre-treatment of low-abundant bio-samples before ELISA, more sensitive and quantitative detection of insulin in serum can be achieved comparing to commercial ELISA kit. This exploratory study suggests a great potential for enhancing the efficiency of current detection strategies by rationally designed nanomaterials.

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Notes and references

- 1 S. Craft, E. Peskind, M. W. Schwartz, G. C. Schellenberg, M. Raskind and D. Porte, Neurology, 1998, 51, 926-926.
- 2 P. H. Sonksen, J Endocrinol, 2001, 170, 13-25.
- 3 F. J. Gil-Bea, M. Solas, A. Solomon, C. Mugueta, B. Winblad, M. Kivipelto, M. J. Ramirez and A. Cedazo-Minguez, J Alzheimers Dis, 2010, 22, 405-413.
- 4 J. Villanueva, J. Philip, D. Entenberg, C. A. Chaparro, M. K. Tanwar, E. C. Holland and P. Tempst, Anal Chem, 2004, 76, 1560-1570.
- 5 M. Thevis, A. Thomas, P. Delahaut, A. Bosseloir and W. Schanzer, Anal Chem, 2005, 77, 3579-3585.
- M. P. Tiwari and B. B. Prasad, J Chromatogr A, 2014, 1337, 6 22-31.
- 7 M. M. Moein, M. Javanbakht and B. Akbari-adergani, Talanta, 2014, 121, 30-36.
- L. Dou and I. S. Krull, Anal Chem, 1990, 62, 2599-2606. 8
- 9 S. Ravi, K. K. Peh, Y. Darwis, B. K. Murthy and T. R. R. Singh, Chromatographia, 2007, 66, 805-809.
- 10 G. Khaksa, K. Nalini, M. Bhat and N. Udupa, Anal Biochem, 1998, 260, 92-95.
- 11 J. Groen, C. E. Kamminga, A. F. Willebrands and J. R. Blickman, J Clin Invest, 1952, 31, 97-106.
- 12 N. T. Ditto, T. R. Kline, P. D. Alfinito and J. R. Slemmon, 2009, 182, 260-265.
- 13 C. Lei, O. Noonan, S. Jambhrunkar, K. Qian, C. Xu, J. Zhang, A. Nouwens and C. Z. Yu, Small, 2014, 10, 2413-2418.
- 14 C. K. Chiang, W. T. Chen and H. T. Chang, Chem Soc Rev, 2011, 40, 1269-1281.
- 15 R. J. Tian, H. Zhang, M. L. Ye, X. G. Jiang, L. H. Hu, X. Li, X. H. Bao and H. F. Zou, Angew Chem Int Edit, 2007, 46, 962-965.
- 16 M. T. Hurley, Z. F. Wang, A. Mahle, D. Rabin, Q. Liu, D. S. English, M. R. Zachariah, D. Stein and P. DeShong, Adv Funct Mater, 2013, 23, 3335-3343.
- 17 S. D. Jayasena, Clin Chem, 1999, 45, 1628-1650.
- 18 M. Holmberg and X. L. Hou, Langmuir, 2009, 25, 2081-2089.
- 19 H. N. Wang, X. F. Zhou, M. H. Yu, Y. H. Wang, L. Han, J. Zhang, P. Yuan, G. Auchterlonie, J. Zou and C. Z. Yu, J Am Chem Soc, 2006, 128, 15992-15993.
- 20 T. G. Cha, B. A. Baker, M. D. Sauffer, J. Salgado, D. Jaroch, J. L. Rickus, D. M. Porterfield and J. H. Choi, Acs Nano, 2011, 5, 4236-4244.
- 21 J. F. Xiao, J. A. Carter, K. A. Frederick and L. B. McGown, J Sep Sci, 2009, 32, 1654-1664.
- 22 X. Y. Zhang, S. C. Zhu, C. H. Deng and X. M. Zhang, Chem Commun, 2012, 48, 2689-2691.

- 23 X. Wang, Y. Liu, S. J. Wang, D. H. Shi, X. G. Zhou, C. Y. Wang, J. Wu, Z. Y. Zeng, Y. J. Li, J. Sun, J. D. Wang, L. J. Zhang, Z. G. Teng and G. M. Lu, Appl Surf Sci, 2015, 332, 308-317.
- 24 H. Salmio and D. Bruhwiler, J Phys Chem C, 2007, 111, 923-929.
- 25 S. M. Darby, M. L. Miller, R. O. Allen and M. LeBeau, J Anal Toxicol, 2001, 25, 8-14.
- 26 W. Kern, C. Benedict, B. Schultes, F. Plohr, A. Moser, J. Born, H. L. Fehm and M. Hallschmid, Diabetologia, 2006, 49, 2790 2792.
- 27 T. Kuuranne, A. Thomas, A. Leinonen, P. Delahaut, A. Bosseloir, W. Schanzer and M. Thevis, Rapid Commun. Mass Spectrom., 2008, 22, 355-362.
- 28 A. Thomas, W. Schanzer, P. Delahaut and M. Thevis, Drug Test Anal, 2009, 1, 219-227.
- 29 S. Pichini, R. Ventura, I. Palmi, S. di Carlo, A. Bacosi, K. Langohr, R. Abellan, J. A. Pascual, R. Pacifici, J. Segura and P Zuccaro, J Pharmaceut Biomed, 2010, 53, 1003-1010.
- 30 E. N. M. Ho, T. S. M. Wan, A. S. Y. Wong, K. K. H. Lam and B. D. Stewart, J Chromatogr A, 2011, 1218, 1139-1146.
- 31 D. H. Pan, P. Yuan, L. Z. Zhao, N. A. Liu, L. Zhou, G. F. Wei, Zhang, Y. C. Ling, Y. Fan, B. Y. Wei, H. Y. Liu, C. Z. Yu and X. J Bao, Chem Mater, 2009, 21, 5413-5425.
- 32 S. Jarnbhrunkar, M. H. Yu, J. Yang, J. Zhang, A. Shrotri, L. Endo-Munoz, J. Moreau, G. Q. Lu and C. Z. Yu, J Am Chem Soc, 2013, 135, 8444-8447.
- 33 M. Kruk, M. Jaroniec, C. H. Ko and R. Ryoo, Chem Mater, 2000, 12, 1961-1968.
- 34 X. S. Zhao and G. Q. Lu, J Phys Chem B, 1998, 102, 1556-1561.
- 35 M. C. Estevez, Y. F. Huang, H. Z. Kang, M. B. O'Donoghue, S Bamrungsap, J. L. Yan, X. L. Chen and W. H. Tan, Methods Mol Biol, 2010, 624, 235-248.
- 36 J. El-Gindi, K. Benson, L. De Cola, H. J. Galla and N. S. Kehr, Angew Chem Int Edit, 2012, 51, 3716-3720.
- 37 J. Buijs, C. C. Vera, E. Ayala, E. Steensma, P. Hakansson and S Oscarsson, Anal Chem, 1999, 71, 3219-3225.
- 38 A. R. Bhat and H. F. Wu, Rapid Commun. Mass Spectrom., 2010, 24, 3547-3552.
- 39 A. Greiderer, M. Rainer, M. Najam-ul-Haq, R. M. Vallant, C. W. Huck and G. K. Bonn, Amino Acids, 2009, 37, 341-348.
- 40 M. Thevis, A. Thomas, P. Delahaut, A. Bosseloir and W. Schanzer, Anal Chem, 2006, 78, 1897-1903.

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