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# Оценка геометрических отклонений, возникающих при воспроизведении трёхмерных моделей средствами аддитивного производства, по данным компьютерной томографии

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#### АННОТАЦИЯ

Обоснование. Технологии трёхмерного моделирования и трёхмерной печати к настоящему времени нашли применение в различных областях клинической и фундаментальной медицины, преимущественно хирургической направленности. Говоря о предоперационной подготовке хирургов, соответствие напечатанных изделий анатомии пациента может играть важную роль в оценке патологических изменений и способах их коррекции. Определение отклонений размеров получаемых моделей сопряжено с этическими и техническими трудностями, связанными с необходимостью определения эталона и проведения большого количества измерений соответственно. В настоящей работе предлагаются использование в качестве эталона геометрической фигуры с заранее известными размерами и оценка линейных отклонений при помощи итеративного алгоритма ближайших точек для каждой из вершин полученной средствами прототипирования полигональной сетки.

**Цель** — оценить геометрические отклонения, возникающие при воспроизведении объектов, имитирующих костную ткань, средствами трёхмерного моделирования (на основе данных компьютерной томографии) и аддитивного производства.

**Материалы и методы.** Для создания исходного объекта использовали программу FreeCAD, редактирование полигональных сеток проводили в программах Blender и Meshmixer. 3D-печать моделей выполняли на принтере Ender-3 из содержащего частицы меди PLA-пластика BFCopper. Сканирование производили 128-срезовым компьютерным томографом Philips Ingenuity CT. Серии томографических изображений загружали в программу 3D Slicer, где на их основе создавали виртуальные модели методами автоматической (с пороговыми значениями 500 HU, 0 HU, -500 HU, -750 HU) и ручной сегментации. Сравнение исходных и воспроизведённых моделей производили на основе итеративного алгоритма ближайших точек в программе CloudCompare.

**Результаты.** В зависимости от метода сегментации объём воспроизведённых моделей превышал объём соответствующих исходных моделей на 1–27%. Средние значения линейных отклонений полигональных сеток воспроизведённых моделей от исходных составили 0,03–0,41 мм. Сравнение значений интегральных сумм линейных отклонений и изменений объёма моделей с использованием коэффициента ранговой корреляции Спирмена показало между ними значимую корреляционную связь (р=0,83; t<sub>амп</sub>=5,27, *p*=0,05).

Заключение. Геометрические параметры воспроизводимого объекта неизбежно изменяются, при этом искажение больше зависит от выбранного способа сегментации, чем от общих масштабов модели или её частей. Использование ручного способа сегментации может привести к большему искажению линейных размеров (по сравнению с автоматическим), но позволяет сохранить все необходимые анатомические структуры.

Ключевые слова: компьютерная томография; 3D-моделирование; 3D-печать; предоперационный период; точность воспроизведения; итеративный алгоритм ближайших точек.

#### Как цитировать

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# Evaluation of geometric deviations in rapid prototyped three-dimensional models created from computed tomography data

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### ABSTRACT

**BACKGROUND:** Computer-aided design and three-dimensional printing have been used in various clinical and fundamental medicine fields, especially in surgery. For example, in the preoperative period, the correspondence of printed products to the anatomy can play an important role in evaluating pathological changes and correction methods. However, determining dimensional deviations of printed models involves ethical and technical difficulties associated with defining a reference and taking many measurements, respectively. Therefore, we propose to use a geometric object with known dimensions as a reference and estimate linear deviations using the Iterative Closest Point algorithm for each of the vertices of the prototyped polygonal mesh.

AIMS: To evaluate the geometric deviations associated with creation of bone-like physical objects from computed tomography data using computer-aided design and additive manufacturing.

**MATERIALS AND METHODS:** The source object was created using the FreeCAD application; Blender and Meshmixer software was used for polygon meshes correction and transformation. The 3D printing was carried out on an Ender-3 printer with copper-impregnated polylactide plastic BFCopper. Scanning was performed using a 128-slice tomograph Philips Ingenuity CT. A series of tomographic images were processed in 3DSlicer software to create virtual models by semiautomatic segmentation with threshold values of 500 HU, 0 HU, –500 HU, –750 HU, and manual segmentation. Reproduced and reference polygon meshes were compared using the Iterative Closest Point algorithm in CloudCompare software.

**RESULTS:** The volume of reproduced models exceeded the volume of respective reference models by 1%–27%. The average point cloud linear deviation values of reproduced models from the reference ones were 0.03–0.41 mm. A significant correlation between integral sums of linear deviations and changes in the volume of reproduced models was shown using Spearman's rank correlation coefficient ( $\rho = 0.83$ ; t<sub>emp</sub> = 5.27, *p*=0.05).

**CONCLUSION:** The geometry of the reproduced object changes inevitably, while the linear deviations depend more on the chosen segmentation method than on the overall size of the model or its structures. The manual segmentation method can lead to greater linear deviations, though it saves all the necessary anatomical structures.

**Keywords:** computed tomography; computer aided design; 3D printing; preoperative period; dimensional measurement accuracy; Iterative Closest Point algorithm.

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# 根据计算机断层扫描数据评估通过增材制造复制三维 模型引起的几何偏差

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### 简评

**论证。**三维建模和三维打印技术现已在临床和基础医学的各个领域得到应用,主要是在外 科领域。谈到外科医生的术前准备,印刷品与患者解剖结构的一致性可以在评估病变和纠正 病变的方法方面发挥重要作用。确定所得模型大小的偏差与伦理和技术困难有关,这些困难 分别与确定标准和进行大量测量的需要相关。 本文中我们建议使用具有预定尺寸的几何图 形作为参考,并使用通过原型制作获得的多边形网格的每个顶点的最近点的迭代算法来估计 线性偏差。

**目标**是通过三维建模(基于计算机断层扫描数据)和增材制造来评估模拟骨组织的物体复制时出现的几何偏差。

材料与方法为了创建初始对象,使用了 FreeCAD 程序,在 Blender 和 Meshmixer 程序 中编辑多边形网格。 这些模型是在含有铜颗粒的 BFCopper PLA 的 Ender-3 打印机上进 行 3D 打印的。 使用飞利浦 Ingenuity CT 128 层 CT 扫描仪进行扫描。 将一系列断层 图像加载到 3D Slicer 程序中,使用自动(阈值为 500 HU、0 HU、-500 HU、-750 HU) 和手动分割的方法创建虚拟模型. 原始模型和复制模型的比较是基于 CloudCompare 程序中 最近点的迭代算法进行的。

**结果。**根据分割方法的不同,复制模型的体积超过相应原始模型的体积 1-27%。 复制模型的多边形网格与原始模型的线性偏差平均值为 0.03-0.41 毫米。 使用斯皮尔曼等级相关系数比较线性偏差和模型体积变化的积分和的值显示它们之间存在显着相关性(ρ = 0.83; t<sub>emp</sub>=5.27, p=0.05)。

**结论。**再现对象的几何参数不可避免地会发生变化,失真更多地取决于所选的分割方法, 而不是模型或其零件的总体比例。 使用手动分割方法会导致线性尺寸的更大失真(与自动 分割方法相比),但它允许您保留所有必要的解剖结构。

关键词:计算机断层扫描; 3D建模; 3D打印; 术前期; 保真度; 迭代最近点算法。

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## BACKGROUND

Medical prototyping technologies, which combine computer-aided design (CAD) and three-dimensional (3D) printing methods, have been increasingly used in various fields of clinical and basic medicine in recent decades [1–4]. Particularly, this was facilitated by the widespread use of additive manufacturing facilities since the 2000s, following the launch of the replicating rapid prototypes and the patent expiration for the fused deposition modeling technology.

The medical prototyping success in the clinical practice depends on the maximum compliance of the resulting products with the patient's anatomy [5]. Thus, in some cases (printing personalized implants and creating guiding resection templates), the degree of congruence between the printed medical product and the bone surface may directly influence the surgical treatment outcome [6]. In other situations (preoperative assessment and creating training phantoms), accurate reproduction of geometric relations between the normal and abnormal tissues predicts possible complications during surgery [7].

Obtaining the reference values for the dimensions of the studied anatomical structures is difficult due to the barrier in the accuracy assessment of the resulting models *in vivo*. Direct measurement of the object of interest is not always possible even during the surgical intervention (due to organ deformations), and obtaining 3D models and assessing their inaccuracies are based on the medical imaging methods used for the noninvasive collection of morphological characteristics of organs. Alternatively, a comparison of models derived from *ex vivo* scans of bone structures may be used; however, this approach uses a relatively small number of checkpoints, and the complexity of their interpretation by different specialists may lead to errors [8]. Tests on 3D models of large mammals (e.g., pigs) give good results with their subsequent introduction into the clinical practice [9].

Physical models were selected as objects of study in earlier works devoted to the accuracy assessment of model reproduction by additive manufacturing tools. These models were measured by several checkpoints using direct means: ruler, caliper, and coordinate measuring machine [10-12]. The present study proposes to use a fundamentally different approach to assess geometric deviations, which consists of all polygonal mesh points comparison obtained using additive technologies with a reference representing the original 3D model. Therefore, a model of complex shape and predetermined dimensions designed using parametric modeling was used as a reference. Conversely, the compared object was a virtual model of a reference passed three main stages of medical prototyping (3D printing, scanning, and modeling). This approach simultaneously assessed the deviations over the entire surface of the product, without the need for numerous linear measurements, while reducing the human factor influence on the measurement process.

This study aimed to assess the geometric deviations that occurred when reproducing objects imitating bone tissues using 3D modeling (based on computed tomography [CT] data) and additive manufacturing.

# MATERIALS AND METHODS

The study design is shown in Fig. 1.



CT: computed tomography.

In the FreeCAD<sup>1</sup> parametric modeling program (FreeCAD Community, Germany), a solid model based on a cuboid with dimensions of  $20 \times 20 \times 12$  mm (l  $\times$  w  $\times$  h) was designed, which contained five end-to-end parallel channels with diameters of 1, 2, 3, 4, and 10 mm. Two recesses and two eminences in the form of a hemisphere and a cone (imitating rounded and sharpened notches and protrusions on the bone surface) at 4 mm in height were created on the upper edge of the model (hereinafter-reference 1). The diameter of the channels was selected to simulate the different foramina of the human skull on one product, which are visualized by modern CT methods. A 1-mm step for smaller diameter channels and irregularities of a given shape on the product surface was applied to manually assess the quality during model printing. Using the Blender<sup>2</sup> software package (Blender Foundation, the Netherlands), volumes were calculated and a copy of this reference, doubled in length, width, and height (hereinafter-reference 2), was created to check the effect of size increase on linear deviations.

Parametric models saved in Standard Triangle Language (STL) format were uploaded to the Repetier Host<sup>3</sup> program (Hot-World GmbH & Co. KG, Germany), where the G-code file for the 3D printer was generated using the CuraEngine<sup>4</sup> slicer (Ultimaker, Netherlands) with the following printing parameters: 0.2 mm layer height, 0.8 mm wall thickness, 33% filling (selected empirically), 50 mm/s speed, 210°C nozzle temperature, 50°C platform temperature, forced model blowing, 5 mm retract, and 100% filament flow. The printing was performed on an Ender-3 3D printer (Creality3D, China) with 0.4 mm nozzle diameter using the BFCopper PLA plastic (Best Filament, Russia) containing copper particles to simulate the X-ray density of bone tissue (average X-ray density of plastic at 100% filling was +1762 HU,  $\sigma = 172$  HU).

The resulting products were scanned using a 128-slice Ingenuity CT scanner (Philips, the Netherlands) in the air with channels oriented perpendicular to the gentry plane. The X-ray tube voltage, current, slice thickness, and pixel size of the reconstructed slices were 120 kV, 117 mA, 0.625 mm, and  $0.43 \times 0.43$  mm, respectively.

Series of tomographic images in Digital Imaging and Communication in Medicine (DICOM) format were loaded into 3D Slicer<sup>5</sup> software (3D Slicer Community, US). The images were used to create STL models of reference 1: four by automatic voxel selection with the threshold values of +500 HU, 0 HU, -500 HU, and -750 HU, respectively (Threshold Paint tool), and one by manual slice voxel tracing (Paint tool). In addition, one model of reference 2 was created by automatic selection of voxels with values > -500 HU. The threshold values for reference 1 were selected empirically based on the X-ray density values in the outer layer of the model (approximately 0.9 mm thick) as -1000...+500 HU (-1000 HU was replaced with -750 HU to exclude the ambient air from the model). Low parameters (-500 HU and -750 HU) were deliberately chosen due to the pronounced defects when using positive values of the segmentation threshold. The density threshold for reference 2 was randomly selected from the thresholds used for reference 1.

The resulting models were loaded into the Meshmixer<sup>6</sup> program (Autodesk, USA), where polygonization errors were analyzed and corrected, structures not in contact with the outer shell of the models were removed, and the model mesh was rebuilt with a fixed polygon edge length (Remesh-Target Edge Length tool) equal to 0.25 mm. The appearance of the model at each of the listed stages is shown in Fig. 2.

At the final stage, the models obtained from CT data, together with their references, were loaded in pairs into



**Fig. 2.** The appearance of the reference model 1 after the following stages: *a*, parametric modeling; *b*, 3D-printing; *c*, CT scanning (axial slice, +805 HU window level, and 3718 HU window width; greater density of protrusions due to closer filament stacking in the horizontal plane), and *d*, creating a polygonal mesh based on CT data. CT: computed tomography.

<sup>&</sup>lt;sup>1</sup> FreeCAD Your own 3D parametric modeler. Available at https://www.freecadweb.org/. Accessed on 05/15/2021.

<sup>&</sup>lt;sup>2</sup> Blender 2.93.2 LTS. Available at https://www.blender.org/. Accessed on 05/15/2021.

<sup>&</sup>lt;sup>3</sup> Repetier. Available at https://www.repetier.com/. Accessed on 05/15/2021.

<sup>&</sup>lt;sup>4</sup> Ultimaker Cura. Available at https://ultimaker.com/software/ultimaker-cura/. Accessed on 05/15/2021.

<sup>&</sup>lt;sup>5</sup> 3D Slicer image computing platform. Available at https://www.slicer.org/. Accessed on 05/15/2021.

<sup>&</sup>lt;sup>6</sup> Autodesk Meshmixer free software for making awesome stuff. Available at https://www.meshmixer.com/. Accessed on 05/15/2021.

CloudCompare<sup>7</sup> software (CloudCompare Project, France), where linear deviations of the final CAD model were calculated based on the iterative closest point (ICP) algorithm. The calculation was performed from each vertex of the obtained polygonal mesh along the normal to the nearest surface of the reference. Statistical analysis was performed using the GNU PSPP<sup>8</sup> program (Free Software Foundation, US).

## RESULTS

Model volumes were determined for a preliminary assessment of their shape distortion. Particularly, if the volume of the rebuilt polygonal mesh was less/more compared to that of the reference CAD model, the linear dimensions of the analyzed model were expected to decrease/increase. In the case of equal volumes, either a size match or a compensated distortion was expected. The volumes of references and models obtained by different methods of segmentation are shown in Table 1.

Polygonal meshes processed with the Remesh tool have an approximately equal density of polygonal vertices distribution per surface unit. When comparing the models (combining cuboidal vertices) with the references, the resulting data set was the number of polygonal mesh nodes removed from the surface of the CAD model at a certain distance (mm) in the outer (positive values) or inner (negative values) directions. An example of aligning the model and the resulting histogram of surface point deviation is shown in Fig. 3. The additional peak in the region of positive values is due to the anisotropy of the voxels along the Z-axis, as well as a slightly excessive material application on the side faces of the cuboid.

Obtained values of distorted geometric dimensions for each model are summarized in Table 2.



Table 1. Volumes of virtual models

Model name, cuboid dimensions, mm (segmentation type and threshold)	Volume, mm <sup>3</sup>	Differences with the reference, mm <sup>3</sup>
Reference 1	3576	-
20×20×12 (auto +500)	3607	31 (0,9%)
20×20×12 (auto 0)	3901	325 (9,1%)
20×20×12 (auto -500)	4255	679 (19%)
20×20×12 (auto -750)	4480	904 (25,3%)
20×20×12 (manual)	4538	962 (26,9%)
Reference 2	28 608	-
40×40×24 (auto -500)	31 140	2532 (8,9%)

The number of measurements performed by the program corresponded to the number of vertices of the polygonal mesh (approximately 100 thousand), and the distribution of deviations of linear dimensions were close to normal. Based on obtained values, the program built a Gaussian (real Gaussian function), with the argument of maximization, which was used as the average deviation of linear dimensions of this model. All linear deviations calculated by the program were divided by their values into intervals of equal width (classes) to build a histogram. For each model, the value of the integral sum (hereinafter—Sum) of linear model deviations was calculated as follows:

$$Sum = \sum_{i=1}^{n} (\mathbf{d}_i \times \mathbf{q}_i), \qquad (1)$$

Where  $d_i$  is the minimum value of the linear deviation in the *i*-th class;  $q_i$  is the number of representatives in the *i*-th class; and *n* is the total number of classes.



Fig. 3. Alignment of polygonal meshes of models (a) and a histogram of the calculated deviations of linear dimensions (b).

<sup>&</sup>lt;sup>7</sup> CloudCompare: 3D point cloud and mesh processing software. Open Source Project. Available at http://www.cloudcompare.org/. Accessed on 05/15/2021.

<sup>&</sup>lt;sup>8</sup> GNU PSPP. Available at https://www.gnu.org/software/pspp/. Accessed on 05/15/2021.

Model name	Minimum value, mm	Maximum value, mm	Average value, mm	Standard deviation, mm	Integral sum, mm
20×20×12 (auto +500)	-0,58	1,146	0,026	0,214	1904
20×20×12 (auto 0)	-0,533	1,019	0,141	0,223	9532
20×20×12 (auto -500)	-0,421	1,129	0,296	0,211	20 756
20×20×12 (auto -750)	-0,675	1,107	0,373	0,197	27 179
20×20×12 (manual)	-0,809	1,068	0,411	0,253	18 190
40×40×24 (auto -500)	-0,862	1,353	0,37	0,275	50 213

Table 2. Linear deviations of model dimensions

Linear normalization by formula (2) was applied to each of the values (sum and volume change) for their bringing to a dimensionless form.

$$\widetilde{\mathbf{X}}_{i} = \frac{\mathbf{X}_{i} - \mathbf{X}_{i, \min}}{\mathbf{X}_{i, \max} - \mathbf{X}_{i, \min}}.$$
(2)

The linearly normalized sum values were compared with the volume change indices of the corresponding models to check the shape distortion direction (Fig. 4).

A comparison of the sums of linear deviations and relative changes in the model volumes using Spearman's rank correlation coefficient showed a high level of correlation ( $\rho = 0.83$ , t<sub>emp</sub> = 5.27, and p = 0.05). Therefore, the change in the reproduced model volume corresponds to the linear deviations measured by the used software tools.

### DISCUSSION

The process of creating a medical prototype includes three main stages: data aquisition (scanning), data processing (creating a virtual model), and 3D printing [13].

The first stage is a radiological examination, which obtains data on the 3D structure of the region of interest with a high spatial resolution (CT or magnetic resonance imaging or 3D ultrasound scanning), results of which are saved as discretized images in DICOM format. The change in geometry at this stage may be due to the specifics of obtaining and processing diagnostic information by the selected imaging method.

The second stage involves segmentation (selecting voxels of medical images related to the created model), voxel mesh into a polygonal mesh conversion, and resulting 3D model editing. Segmentation may be of three types: manual (fully performed by the operator), semiautomatic (performed by the computer and corrected by the operator), and automatic (fully performed by the computer) [14]. Automated methods are more attractive due to lower labor costs; however, because of characteristics of used computer algorithms, the geometry of the final product may be severely distorted, and their accuracy requires a separate study [15]. At this stage,



Fig. 4. Linear normalized values. Red color: differences in the volume of models with the reference; blue color: differences in the integral sum of linear deviations.

a specialist with knowledge in anatomy (particularly, radiological anatomy) should be involved to correctly select and edit the objects of interest. Thus, geometry distortion at this stage mainly results from the human factor.

The third stage is the 3D printing of the resulting virtual model. For its successful implementation, the model is pre-processed (slicing) and post-processed (removal of supports and surface handling) after printing (if necessary). The contribution of this stage to the distortion of the final product geometry primarily depends on the type of used additive equipment.

Therefore, to assess geometric distortions in the prototyping process, all three stages must be performed on some physical phantom with known linear dimensions and defined design elements. Scanning conditions and printing settings may affect the accuracy of the resulting product [10, 16]. Thus, the relevant parameters were kept at the same level for all produced models.

Many different software packages can be used to segment medical images. The use of 3D Slicer at the stage of virtual model creation from the DICOM data was due to its accessibility (distributed as open-source software) and a large number of additional modules and extensions that make it an ideal tool for preoperative planning [17]. The results confirm the "dumbbell" effect described earlier, according to which a decreased automatic segmentation threshold leads to an expansion of resulting model outlines [8]. Thus, when the threshold was decreased from +500 to -750 HU, the average value of the surface point displacement from the reference consistently increased from 0.026 to 0.373 mm, respectively. These distortions depend more on the segmentation method compared to the size of the scanned object, since the average linear deviations were 24% higher in all dimensions when the model was proportionally doubled compared to the original size models with the same threshold, and 10% lower as opposed to the original size of manually segmented models.

The relatively high values of geometric deviations in the manually segmented model (by 0.41 mm for a  $20 \times 20$  mm sample) may result from the operator's aligning the side faces of the cuboid along the edges, which were somewhat displaced outward (during 3D printing). Moreover, similar deviations were observed during semiautomatic segmentation. Remarkably, these models reproduced the original shape better due to the preservation of all control elements and the absence of wall defects (Fig. 5).

The visual assessment of models that are automatically segmented with a threshold of 0 and +500 HU observed



**Fig. 5.** The appearance of models segmented semi-automatically with a cutoff threshold of 0 HU (*a*, using a map of deviations from the reference; *b*, general view) and manually (*c*, using a map of deviations from the reference; *d*, general view).



Fig. 6. Measurement of the linear deviations from the reference (blue lines) in the defect area of the compared model (red lines).

1.2 mm thick wall defects, whereas models with -750, -500, and 0 HU thresholds lacked created end-to-end holes with a diameter of 1 mm. The defects with a width exceeding the module of minimum geometry deviations are explained by the thickness measurements performed from two surfaces, each with appropriate deviations (Fig. 6). Thus, the linear displacement summation of the vertex relative to the outer surface ( $d_1$  segment) with the nearby vertex displacement relative to the inner surface ( $d_2$  segment) may correspond to a wall defect of l thickness exceeding the maximum linear deviation value. Similarly, the positive deviation summation leads to the "obliteration" of holes that exceed the maximum size deviation calculated from the ICP algorithm results. Thus, the linear dimensions of the reproduced models include two values of linear displacement.

In addition, channels with a diameter of 1 mm, which were not detected on the mentioned copies during the visual assessment, were not evaluated by the algorithm since the polygonal vertices corresponding to these channels were absent on the models.

### Study limitations

The limitations of this study could be the relatively small sample size and the use of materials that did not correspond to the true bone tissue composition.

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### CONCLUSIONS

Distortion of the original shape inevitably occurs in the creation of a product using medical prototyping (CT scanning, 3D modeling, and 3D printing). The change in the model volumes and the average linear deviations of their surface points from the reference, determined using the ICP algorithm, have positive values. The distortion depends more on the selected segmentation method compared to the overall scale of the model or its parts. The use of the manual seqmentation compared to the semiautomatic method leads to a slightly higher distortion of linear dimensions resulting from aligning the model to the unreliable landmarks; however, it reproduces all elements of the original sample. Thus, for necessary visualized anatomical structure preservation, the resulting virtual models should be corrected by a specialist. In this case, the use of different thresholds in areas with dense and sparse structures may be sufficient for segmentation. Using the semiautomatic segmentation is more preferable if several small structures can be neglected and the main task is to match the overall product dimensions to the simulated organ; however, the thresholds of this method should be selected experimentally depending on the tissue composition and scanning parameters.

# ADDITIONAL INFORMATION

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