with the same frequency. Patterns 1 and 2 have larger contact surface for the talus compared to pattern 3. Male bones have larger contact surface for the talus than female bones.

Conclusions. The sum of pattern 1 and pattern 3 frequencies in our study was high, so we concluded that our population is more prone to subtalar arthritis. Knowing how frequent are the different patterns of calcaneal facets for the talus in a certain population is important for orthopaedic surgeons when performing foot osteotomy.

THE PECULIARITIES OF HUMAN AND ANIMAL HUMERUS SYSTEMIC ORGANIZATION AT THE DISTAL EPIPHYSIS LEVEL

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Aim. The main purpose of research is to detect the general patterns and characteristics of the Humerus (H) systemic organization (SO) of the man's and animals with different volumes of movements in the elbow joint at the level of it's distal epiphysis (DE).

Material and Methods. The H Bos tautus Taurus (B) R40/L40, Orycolagus cuniculus (O) — R40/L40, Canis lupus familiaris (C) — 39R/39L, Homo sapiens (H) — R77/L77 were studied using osteometry. There are 23 homologous structures were measured on each bone. A group of angular structures of the DE: corner of torsion, tilt of the diaphysis, the angle of groove of the block up to the plane of condyles. All values were analyzed involved calculating confidence intervals at $\alpha \ge 0.05$ and multilevel Maximum likelihood factors Analysis (FA) with the rotation Equamax normalized and taking into account the lateralization of bones with the calculation of factor loads (FL).

Results and Discussion. There were no differences in the magnitude of the R/L angles. The value of the twist angle varies in the O-B-H-C series from 46 to 81°. The angle of inclination of the diaphysis increases in the series H-B-O-C is from 6 to 37°. Angle of the groove of the block, representing the rotation of the lower limb-O-C-B-H is in the range $46-59^{\circ}$.

Conclusions. The results of the study revealed the presence of a latent asymmetry of the SO of the H at the level of it's DE, which is caused by a change in the number and volume of movements in evolution. STUDY OF MYOCARDIAL REMODELING MARKERS IN EXPERIMENTAL HISTOTOXIC HYPOXIA

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Key words: myocardium, rats, fibrogenesis, hypoxia

Background. In modern works, there are isolated data concerning the problem of cardiac remodeling and fibrogenesis.

Aim. The available data are ambiguous, so the goal of the study is to study the processes of fibrogenesis and inflammation in the myocardium of sexually mature rats.

Material and Methods. The study was conducted on 25 mature male Wistar rats weighing 220– 310 g. The control group was represented by 5 intact animals. A solution of cobalt chloride for the modeling of histoxic hypoxia was introduced through an atraumatic probe into the stomach at a dose of 4 mg/ kg, daily once a day for 30 days.

Results and Discussion. The conducted study showed that changes in the myocardium developed, which, according to morpho-functional manifestations, could be attributed to varieties of toxic cardiomyopathy. Hydrophobic dystrophy was observed in the cytoplasm of cardiomyocytes, myocytolysis foci appeared. Masson's staining revealed an excessive amount of collagen fibers in the interstitium, and also in the perivascular space, which apparently made it more difficult to transport substrates and oxygen from the bloodstream to the working cells under hypoxic conditions. Perivascularly and in interstitium, intensely colored macrophages and fibroblasts appeared. Remodeling of the myocardium, which occurs during hypoxic damage, is accompanied by a reorganization of the connective tissue stroma. At the same time, the enzymatic cleavage of the previously existing connective tissue fibers develops and de novo synthesis of interstitial substance according to the new conditions of mechanical loads. The leading role in this process is assigned to the enzymes of the metalloproteinase group. Expression of MMP-9 metalloproteinase was characterized as weakly positive (+) and positive (++) in some parts of the myocardium. Increased expression