

Generation of expanded primary hepatocytes (upcyte® cells) in large quantities for for cell based toxicity and metabolism screenings

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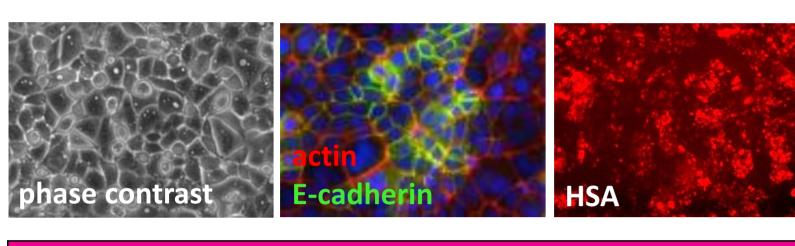
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INTRODUCTION

Isolated liver cells, such as hepatocytes, liver sinusoidal endothelial cells (LSECs), Kupffer Cells and hepatic stellate cells, are frequently used to study hepatic metabolism, toxicity and disease pathogenesis. However, the current in vitro models exhibit several disadvantages, e.g. short culture longevity and artificial culture conditions focusing on a single cell type in 2D culture. The use of primary cells in vitro is compromised by the limited quantity of cells that can be isolated from one donor, a lack of or very restricted proliferation capacity (hepatocytes and LSECs) and/or the change from a quiescent to an activated state (hepatic stellate cells). To overcome this, we have developed a technique which causes primary human hepatocytes to proliferate up to 40 population doublings whilst retaining an adult and metabolically competent phenotype with phase I (Cytochrome P450) and phase II activities when cultured at confluence. The resulting cells are called "upcyte® hepatocytes" and combine proliferation with drug metabolizing activity, a feature which makes them uniquely applicable to metabolism and toxicity studies.

RESULTS

metabolism & characteristics



Specific activity (pmol/min/mg protein)				
Cells	CYP1A2	СҮР2В6	CYP2C9	СҮРЗА4
#10-03	3.3 ± 0.4	40.3 ± 6.5	91.8 ± 5.5	21.4 ± 9.6
#151-03	0.7 ± 1.4	71.1 ± 11.3	29.1 ± 21.4	77.8 ± 22.6
#422A-03	2.3 ± 0.1	33.6 ± 11.4	4.8 ± 3.1	42.9 ± 6.3
#653-03*	17.1 ± 0.5	68.4 ± 18.4	16.2 ± 0.9	178.3 ± 17.0
HepaRG	10.0 ± 1.5	6.45 ± 0.97	4.57 ± 2.93	48.5 ± 13.9
*Donor 653-03-2D6 available:				

upcyte[®] hepatocytes display an adult phenotype

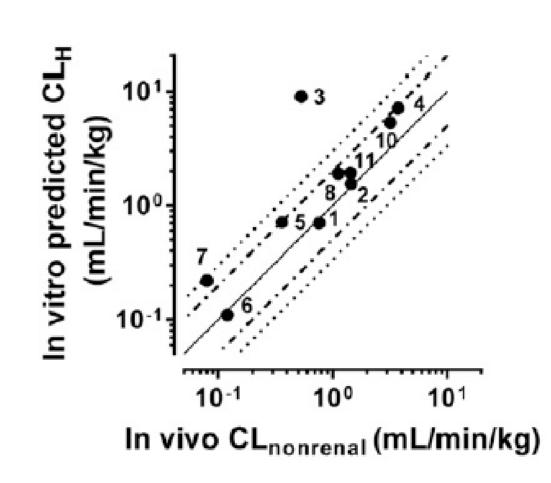
hepatocytes upcyte® express characteristic adult marker proteins (CK8, CK18, HSA, AAT), but lack embryonic markers such as AFP. They further show capability for glycogen storage, urea secretion and albumin synthesis (data not shown).

The cells exhibit basal activity of CYP1A2, 2B6, 2C9, 3A4 and other CYPs in a donor-dependent manner. All enzymes can be induced an inhibited.

upcyte® hepatocytes have similar phase II activities compared to primary hepatocytes

Major hepatic phase II enzymes in humans are UDP-glucuronosyltransferase (UGT), sulfotransferase (SULT) and glutathione S-transferase (GST). Phase II enzyme activities in upcyte® hepatocytes generated from different donors were similar to those of freshly isolated primary human hepatocytes (data not shown).

clearance prediction



Data represent mean 6±S.D. of triplicate incubations per compound (n = 3) determined at day 7 in sandwich culture. Solid line represents conformity, and dashed lines two- and threefold error range. The set of reference drugs was subdivided into low and intermediate cleared compounds (shown here: low): alprazolam (1), prednisolone (2), diazepam (3), voriconazole (4), tolbutamide (5), meloxicam (6), warfarin (7), glimepiride (8), riluzole (10), oxazepam (11).

Boehringer Ingelheim Pharma GmbH & Co. KG, DMPK, Biberach an der Riss, Germany

upcyte® hepatocytes are a potent in vitro tool for the prediction of hepatic clearance (CL_H)

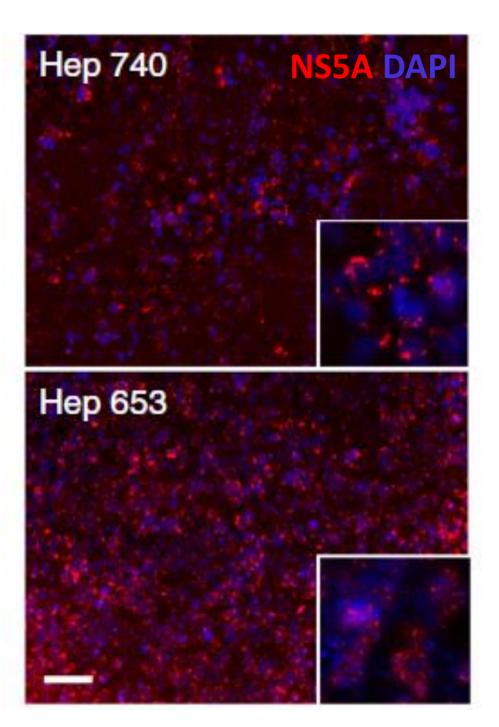
Correlation between in vitro predicted and in vivo CL_{nonrenal} was demonstrated using the well-stirred model disregarding plasma protein binding for low and intermediate clearance compounds (donor 151-03).

Good correlation between predicted CL_H and observed in vivo CL values was observed for the subset of low CL drugs (shown here). CL_H for 73% (8 of 11 compounds) were predicted within twofold of in vivo CL_{nonrenal} and within threefold for 82% (9 of 11 compounds) with a trend for overpredicting the actual in vivo rate.

Schaefer et al., (2015) - Drug Metabolism & Disposition upcyte® human hepatocytes: a potent in vitro tool for the prediction of hepatic clearance of metabolically stable compounds

viral infection: HCV

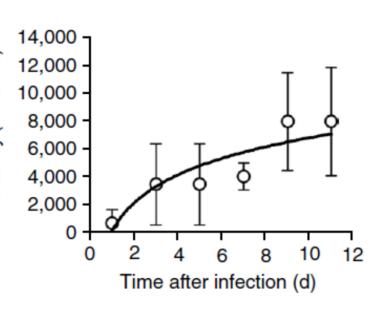
CYP2D6 recombinantly expressed, specific activity for 2D6 1627.5 ± 22.4



upcyte® hepatocytes support HCVcc infection

HCV infects hepatocytes in vivo, but infectivity of primary human hepatocytes in vitro is minimal. To assess whether our upcyte® hepatocytes support the full lifecycle of the HCV cell culture variant **HCVcc**, we exposed differentiated hepatocytes (donors 740 and 653-03) to culture medium containing infectious particles of the JC1 genotype expressing an NS5A-RFP fusion protein. Hepatocytes from both donors showed strong NS5A staining. More than 80% of the cells were infected in both cultures.

levels significantly NS5A-RFP increased over time, reaching a 20-26-fold induction at day 9 post infection. **Production of infectious** particles showed a similar trend, stabilizing after day 9 and reaching 8,000 focus-forming unit (FFU) per milliliter for donor 653-03.



Levy et al., (2015) – Nature Biotechnology long term culture and expansion of primary human hepatocytes The Hebrew University of Jerusalem, Israel / upcyte technologies GmbH

CD31 (protein)

<u>LSECs</u> (brown

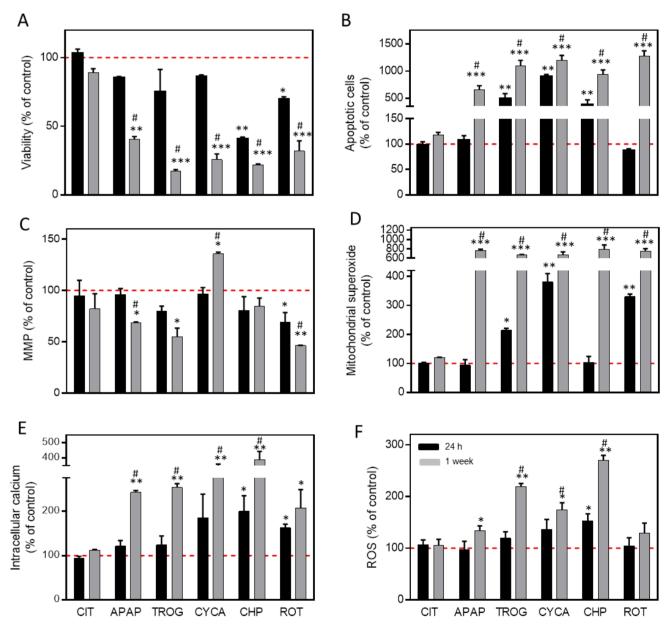
inner ring of

"blossom"

cytotoxicity - acute and repeated-dose studies

sub-cytotoxic concentrations

Exposure time length had dramatic effects on the toxicity profile of a compound. For APAP, no effect was observed after 24 h, whereas 1-week significantly induced treatment apoptosis, mitochondrial depolarization, ROS production and intracellular Ca²⁺ levels. Other tested compounds caused some effects after 24 h, although a significant difference was detected between the two incubation periods at the lowest concentration. CIT (non hepatotoxic control) did not produce any significant effects after 24 h or 1 week treatment.

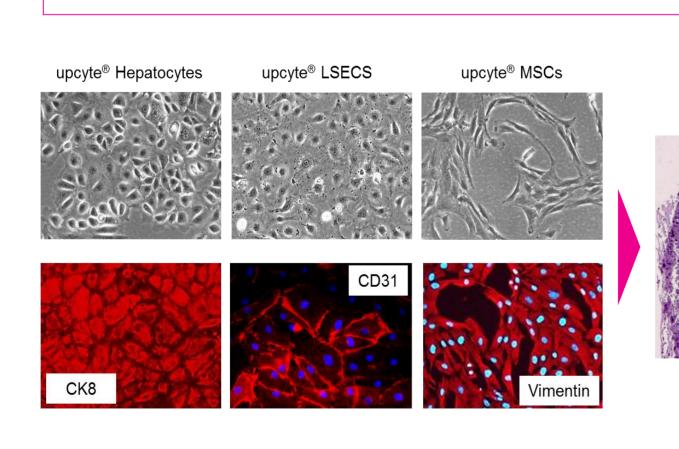


upcyte® hepatocytes (422a-03) were exposed to test compounds for 24 h or 1 week. Fluorescent probes were subsequently employed to evaluate (A) viability, (B) apoptosis, (C) changes in mitochondrial membrane potential (MMP), (D) production of mitochondrial superoxide, (E) ROS (F) intracellular Ca^{2+} levels using HCS. Data are expressed as mean \pm SEM as percentages normalized on untreated control cells. Statistical analysis was performed using Student t-test (*p<0.05, **p<0.01, ***p< 0.001 vs. untreated; #p<0.05 vs. 24 h).

Tolosa et al., (2016) – Toxicological Sciences human upcyte® hepatocytes: characterization of the hepatic phenotype and evaluation for acute and long-term hepatotoxicity routine testing - University of Valencia, Spain

genotoxicity: micronuclei (MN) assay

3D format and co-cultures



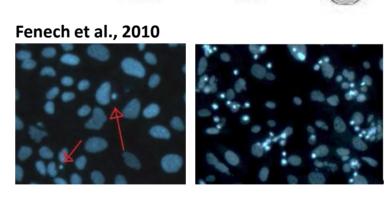
HIF-1α (RNA)

Ramachandran et al., (2016) - Plos One

self-assembly in MatrigelTMcoated 24-well plates

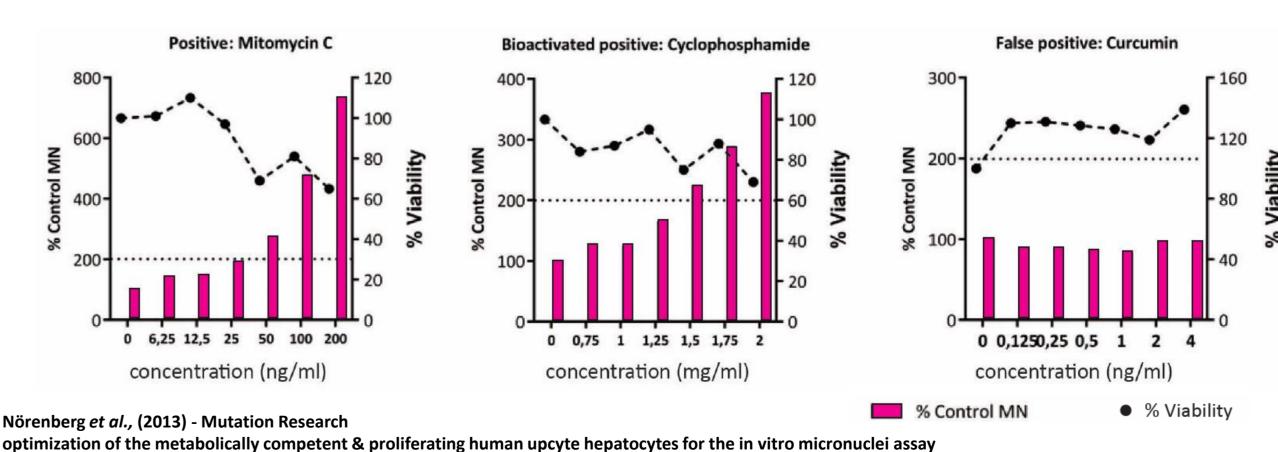
After 72h, cells showed formation of structures resembling liver blossom. Hepatocytes were located in the center and formed the outer petals (HIF- 1α). **LSECs** formed an inner ring (CD31), whereas **HSCs** revealed a typical, "star-shaped" morphology (α -SMC).

αSMA (protein) morphology)



upcyte® hepatocytes can discriminate compounds of 3 classes

The requirements for the MN assay are proliferating and metabolically competent cells. While pHH lack proliferation, cells lines lack expression of metabolizing enzymes. The use of S9 extract as exogenous metabolizing system involves a high "false positive" rate. Incubation for 96 h (long term treatment for 2 population doublings) allowed upcyte® cells to discriminate correctly between true positives (direct and bio-activated), true negatives and false positives.



SUMMARY & CONCLUSION

In vitro generation of functional liver organoid-like structures using adult human cells

"blossom"

In conclusion, upcyte® hepatocyte cultures are characterized by a differentiated phenotype and exhibit functional phase I, phase II and transporter activities. These data support the use of upcyte® hepatocytes for various applications, such as as metabolism & toxicity screening assays, viral infection and 3D culture. Moreover, this technology allows for the generation of large batches of upcyte[®] hepatocytes (up to 12 x 10⁹ cells per donor), enabling a reproducible and standardized experimental setting.

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