

Evaluation and Synthesis of Polar Aryl- and Heteroaryl Spiroazetidone-Piperidine Acetamides as Ghrelin Inverse Agonists

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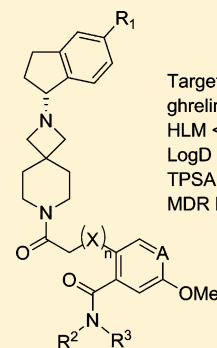
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Supporting Information

ABSTRACT: Several polar heteroaromatic acetic acids and their piperidine amides were synthesized and evaluated as ghrelin or type 1a growth hormone secretagogue receptor (GHS-R1a) inverse agonists. Efforts to improve pharmacokinetic and safety profile was achieved by modulating physicochemical properties and, more specifically, emphasizing increased polarity of our chemical series. *ortho*-Carboxamide containing compounds provided optimal physicochemical, pharmacologic, and safety profile. pH-dependent chemical stability was also assessed with our series.



KEYWORDS: Heteroaryl acetic acid, carboxamide, spiroazetidone-piperidine, growth hormone secretagogue receptor 1a, ghrelin, type 2 diabetes

Type 2 diabetes mellitus (T2DM) is the more common form of diabetes and millions of people are affected by T2DM, especially in Western countries.¹ The pancreas of type 2 diabetics cannot either produce insulin or the cells in the body are incapable of using insulin effectively. In addition, about 90% of type 2 diabetics are overweight or obese.² Ghrelin hormone, an acylated 28-amino acid peptide, is a physiological regulator of insulin release and glucose homeostasis,¹ and it plays a central role as a hunger hormone.³ Ghrelin is an endogenous ligand of type 1a growth hormone secretagogue receptor (GHS-R1a), also called GRLN (human ghrelin receptor), which is expressed in the pancreatic islets, and is

released in the pancreatic microcirculation.⁴ Small molecule agonists of ghrelin have been shown to induce hyperglycemia and insulin resistance,^{5,6} whereas antagonists (and inverse agonists) have been shown to improve glucose tolerance, suppress appetite, and promote weight loss.⁷⁻¹⁸

We have previously disclosed our efforts to identify centrally and peripherally acting inverse agonists of ghrelin from the

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spiropiperidine-azetidine series.^{19–21} Our discovery efforts focused on the identification of inverse agonists over antagonists in order to reduce basal receptor firing, as there is a high constitutive activity of ghrelin receptor.²² Our centrally acting and orally bioavailable lead compound **2** (Figure 1)

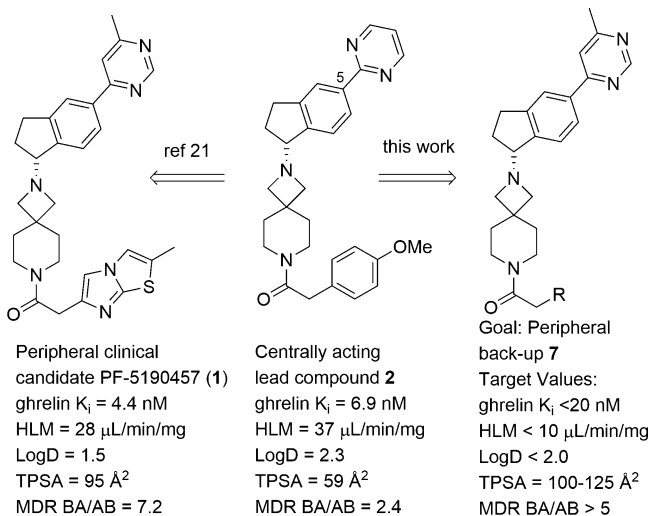
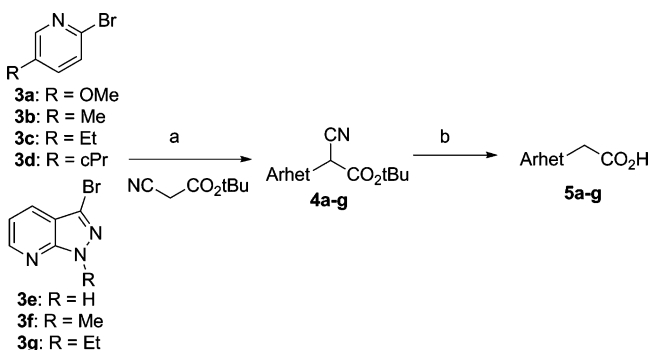


Figure 1. Pfizer ghrelin inverse agonists.

Scheme 1^a

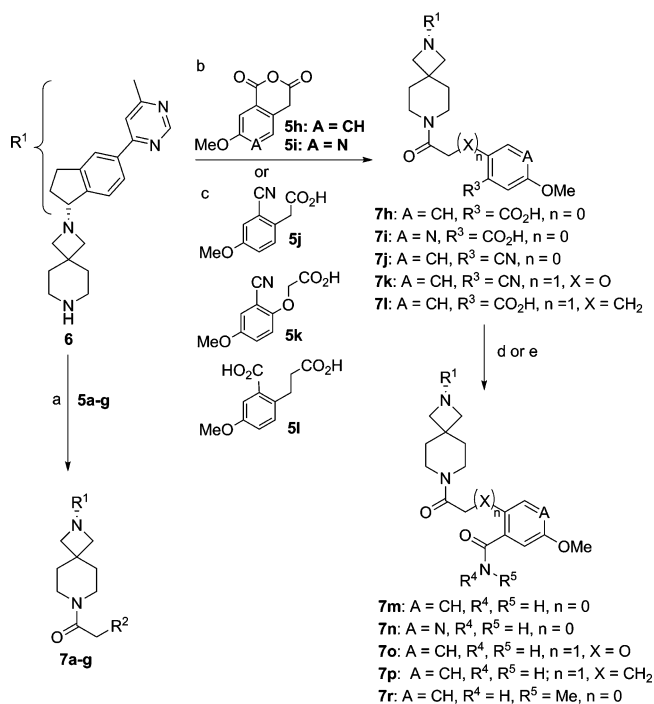


^aReagents and conditions: (a) $\text{Pd}(\text{PPh}_3)_4$ or $\text{Pd}(\text{dppf})_2\text{Cl}_2$, THF or 1,4-dioxane, reflux, 18 h; (b) conc. HCl, reflux, 6–18 h.

showed significant glucose-dependent insulin secretion in a human dispersed islet cell assay. However, compound **2** suffered from several off-target activities in a Cerep safety panel, and the physicochemical properties were not ideal for further development.²⁰ The project team noted that the requisite effect should be derived from action on the islet cells in pancreas; as such, peripheral restriction will have advantages in safety and side effect profile.²⁰

Variations of **2** at both the C(5) position of the indane ring and the piperidine acetamide portion to access compounds of increased lipophilic efficiency (LipE)^{23,24} were investigated and compound **1** (PF-5190457) was discovered as a peripherally acting potent ghrelin inverse agonist.²¹ This compound had excellent off-target selectivity and good predicted human pharmacokinetic properties (projected 35 mg bid dose), and it was advanced to human clinical studies. Concurrently, an alternative strategy was investigated to increase polarity and TPSA of the lead compound **2** with further variations at the acetamide portion. Herein, we describe efforts to identify an improved derivative of compound of **2** with reduced off-target

Scheme 2^a



^aReagents and conditions: (a) EDCl, HBTU or CDI, DIEA or Et₃N; (b) Et₃N; (c) EDCl or CDI, DIEA or Et₃N; (d) Pt(PMe₂OH)₃H, water, EtOH; (e) CDMT·NMM, NH₄Cl or MeNH₂.

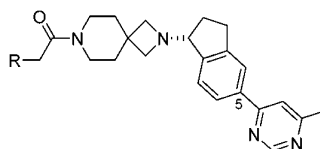
pharmacology, optimal physicochemical properties, and increased polarity.

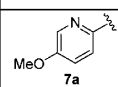
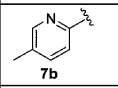
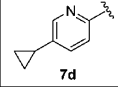
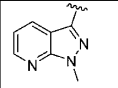
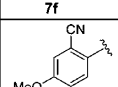
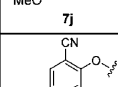
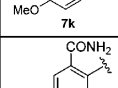
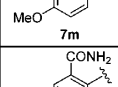
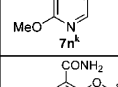
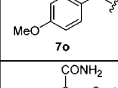
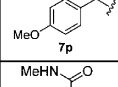
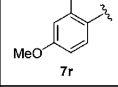
To achieve the desired improvements, we set our goals to achieve at least 20 nM ghrelin potency, lower than 10 $\mu\text{L}/\text{min}/\text{mg}$ intrinsic clearance in human liver microsomes (HLM), logD lower than 2, total polar surface area (TPSA) between 100 and 125 \AA^2 , and efflux ratio (BA to AB) greater than 5 in the PgP efflux assay in MDR-MDCK1 (MDR) cell line.²⁵ The latter two goals were set to minimize brain exposure.^{26,27} With the discovery of **1**, we had optimized the indane C(5) heteroaryl group as the 6-methyl-4-pyrimidine and chose to investigate polar replacements of the 4-methoxyphenyl group of **2**.

The synthesis of several polar heteroaryl acetic acids is shown in Scheme 1 starting from commercially available heteroaryl bromides **3**. Treatment of **3** in the presence of palladium tetrakis(triphenylphosphine) ($\text{Pd}(\text{PPh}_3)_4$) or palladium 1,1'-bis(diphenylphosphino)ferrocene dichloride ($\text{Pd}(\text{dppf})_2\text{Cl}_2$) and an anion generated from *tert*-butylcyanomalonate and KO*t*Bu (or LiO*t*Bu) provided intermediates **4**. Acidic hydrolysis and decarboxylation provided acids **5**. It is worth noting that a second decarboxylation of the final compounds **5b** and **5c** occurs very rapidly (1–2 h) at room temperature and over time (~2 weeks) in a refrigerator. Decarboxylation of similar substrates under basic conditions and heat are known.²⁸ The cyclopropyl substituted acid **5d** was slightly more stable than the methyl and ethyl substituted acids **5b** and **5c**, but even **5d** underwent decarboxylation over 1–2 months at +4 °C. These three acids were freshly prepared and used immediately in the subsequent amide coupling reactions. The other acids (**5a,e,g**) were stable at ambient temperature.

The acids **5a–g** were reacted with amine **6**^{21,29,30} in the presence of a coupling reagent (such as HBTU, TBTU, EDCl,

Table 1. In Vitro Data for Selected Ghrelin Inverse Agonists 7a–s



R	SFLogD ^a	ghrelin binding K _i (nM) ^{b,c}	ghrelin functional K _b (nM) ^{c,d} (%effect) ^e	HLM (μL/min/mg) ^f	TPSA (Å ²) ^g	MDR (BA/AB) ^h	Passive permeability (10 ⁶ cm/s) ⁱ	Stable at pH 7.4 (yes/no) ^j
 7a	1.6	14±2	17±3 (-25%)	19	71.4	6.4	13	yes
 7b	(1.1)	49±4	N/A	20	62.2	4.5	10	yes
 7d	(1.8)	6.4±0.4	14±7 (-15%)	48	62.2	N/A	10	yes
 7f	2.1	166±25	N/A	32	80.4	6.9	12	yes
 7j	2.4	13±2	28±8 (-16%)	20	82.3	1.16	14	yes
 7k	2.3	4.0±0.6	5.0 (n=1) (-14%)	39	91.6	3.9	5.0	yes
 7m	1.4	9.2±4.0	14±3 (-18%)	<8.3	101.6	>17	2.2	no
 7n ^k	(0.8)	263±6	328 (n=1) (-15%)	N/A	114.5	N/A	N/A	no
 7o	2.0	127±17	81 (n=1) (-12%)	11	110.9	>17	4.0	yes
 7p	1.9	139±24	95 (n=1) (-13%)	71.7	101.6	N/A	3.1	yes
 7r	(0.9)	20±1	N/A	13	87.7	>17	13	no
 7s	1.6	28±4	57±18 (-16%)	33.0	87.7	12.0	15	yes

^aSFLogD, shake-flask logD at pH 7.4; if not measured, calculated (ACD Laboratories program v12) LogD at pH 7.4 is given in parentheses. ^bAll human ghrelin receptor SPA binding data were generated as previously described in ref 19. ^cGeometric mean ± standard error of ≥3 measurements, unless noted otherwise. ^dhuman ghrelin receptor agonist/antagonist/inverse agonist GTP-γ-S functional assay as published in ref 20. ^eNegative % effect indicates an inverse agonist. ^fHLM, human liver microsome clearance; not corrected to fraction unbound. ^gTPSA, topological polar surface area. ^hMDR efflux ratio (BA/AB) in MDCK/MDRI cell line as described in ref 25. ⁱPassive permeability measured using low efflux MDCKII cells as described in ref 34. ^jUPLC-MS, ultra pressure liquid chromatography mass spectrometry determination of disappearance of the parent compound and appearance of compound 6 at pH 7.4 over 24 h time period; “yes” indicating stability and “no” indicating instability (see Supporting Information). ^kCompound 7n has indane C(5) 4-methyl-2-pyrimidine substituent (see Supporting Information) instead of 6-methyl-2-pyrimidine substituent that all other analogues have.

or CDI) to form amides 7a–g that were tested in the human ghrelin binding assay,¹⁹ in HLM and MDR assays (Scheme 2, arrow a; and Table 1). Even though compounds 7a and 7d were very potent ghrelin binders (14 and 6 nM, respectively)

and inverse agonists in the functional assay (17 and 14 nM, respectively),²¹ they either did not achieve the required stability in HLM (48 μL/min/mg for 7d) or they suffered from off-target activities (i.e., >50% inhibition of M1, M2, MCH1, and

Table 2. Comparison of Physical Chemical, Pharmacological, ADME, and Safety Properties of Compounds 7m and 2

property	compd 7m	compd 2
MW	525.6	468.6
cLog P	1.76	2.82
SFLogD (pH 7.4) ^a	1.37	2.30
TPSA (Å ²)	102	59
binding K _i (nM) ^{b,c}	9 ± 4	7 ± 2
functional K _b (nM) (% effect) ^{d,e}	14 ± 3 (-18%)	17 ± 7 (-39%)
RLM Cl _{int} (μL/min/mg) ^f	77.4	55.4
DLM Cl _{int} (μL/min/mg) ^g	16.8	29.0
HLM Cl _{int} (μL/min/mg) ^h	<8.6	36.8
CYP450 IC ₅₀ (1A2, 2C9, 2C19, 2D6, 3A4)	all >30 μM	3A4: 2.5 μM
kin sol (pH 6.5, μg/mL) ⁱ	185	187
permeability AB (P _{app} cm/s) ^j	2.2 × 10 ⁻⁶	13.2 × 10 ⁻⁶
permeability (MDCK/MDR1, BA/AB) ^k	>17	2.4
hERG (patch clamp) IC ₅₀ (μM)	47	2.9
CEREP panel (82 assays) IC ₅₀	>10 μM (81 assays)	several hits at 10 μM (ref 21)
CEREP hit (5-HT _{2B}) IC ₅₀ (μM)	7.6	4.3

^aSFLogD, shake-flask logD at pH 7.4. ^bGeometric mean ± standard error of ≥3 measurements. ^cHuman ghrelin receptor SPA binding assay as published in ref 19. ^dHuman ghrelin receptor inverse agonist GTP-γ-S functional assay as published in ref 20. ^eNegative % effect indicates an inverse agonist. ^fRat liver microsome (RLM) intrinsic clearance. ^gDog liver microsome (DLM) intrinsic clearance. ^hHuman liver microsome (HLM) intrinsic clearance. ⁱAnaliza kinetic solubility (kin sol) from 10 mM DMSO solution at pH 6.5 (2% sodium phosphate buffer) after 24 h. ^jPassive permeability using low efflux MDCKII cells as described in ref 34. ^kMDR efflux ratio (BA/AB) in MDCK/MDR1 cell line as described in ref 25.

5HT_{2B} receptors at 10 μM concentration in the Cerep panel for 7a). Compounds 7b–c and 7e–g did not pass our potency or clearance requirements (all >50 nM in the ghrelin binding assay).

An alternative approach to improve clearance via further increases in polarity (logD and TPSA) involved installing a polar substituent, such as a carboxamide or a nitrile, to the aromatic ring of 2. Synthetic routes to these polar amides are depicted in Scheme 2 (arrows b–e). Direct coupling of CN-aryl acids 5j and 5k with amine 6 under EDCI or CDI amide coupling conditions delivered compounds 7j and 7k that could be further transformed to the carboxamides via Pt catalyzed hydrolysis³¹ or via urea-hydrogen peroxide mediated hydrolysis³² to provide compounds 7m and 7o in quantities that were suitable for structure–activity relationship (SAR) purposes.

Compounds 7j, 7m, and 7r (Table 1) exhibited the required target values for ghrelin potency (13, 9.2, and 20 nM,

respectively), low clearance (20, <8.3, and 13 μL/min/mg, respectively), high polarity, and high efflux ratios (>17 MDR BA/AB for 7m and 7r). Compound 7j was among the most potent, but suffered from low MDR BA/AB ratio (1:1). As compound 7m started to emerge as a potential back-up compound, more material needed to be synthesized. The overall yield via the nitrile intermediate 7j was low and not suitable for scale up work. An alternative way to generate the *ortho*-amides was to treat the amine 6 with anhydrides 5h or 5i or *ortho*-acid 5l to generate *ortho*-carboxylic acid intermediates 7h, 7i, and 7l that were then treated with NH₄OH in the presence of 2-chloro-4,6-dimethoxy-1,3,5-triazine (CDMT)³³ and *N*-methylmorpholine (NMM) to give 7m, 7n, and 7p in high and scalable (7m up to 10 g) yields. The intermediate 7h could also be treated under the CDMT conditions in the presence of methylamine to access 7r or in the presence of dimethylamine to access the dimethyl analogue (R₄ = R₅ = Me) in high yields and high purity.

Compound 7m exhibited the optimal combination of potency/clearance/polarity, and it was profiled further in ADME, safety assays, and in vivo pharmacokinetic experiments (Tables 2 and 3). Significant improvements compared to our lead compound 2 were achieved (Table 2). Compound 7m had relatively low microsomal clearance in dog and human liver microsomes (16.8 and <8.6 μL/min/mg, respectively) and elevated clearance in rat liver microsomes (77.4 μL/min/mg). It did not inhibit any of the cytochrome P450 enzymes at 3 μM concentration, and the major clearance mechanism was CYP3A4/3A5 mediated. Compound 7m had very little off-target activity in the Cerep panel and only hit 1 out of 82 targets at 10 μM concentration. The receptor that was inhibited by 7m was 5-HT_{2B} with an IC₅₀ of 7.6 μM. Compound 7m had an MDR-MDCKI BA/AB ratio of higher than 17 indicating potential for impairment through the blood–brain barrier. Passive permeability was relatively low as judged from the permeability value of 2.2 × 10⁻⁶ cm/s (passive permeability (P_{app}) measured using low-efflux MDCKII cells).³⁴ Pharmacokinetic data of 7m was collected in three preclinical species (rat, dog, and monkey; Table 3). Good oral bioavailability was observed in dogs and monkeys (67% and 32%, respectively). It is worth noting that despite high clearance observed in rats, a reasonable oral bioavailability was achieved (15%). This is likely due to “flip-flop” kinetics where the rate of absorption is much slower than disposition.³⁵ Compound 7m had no liabilities in the transformed human liver epithelial (THLE > 248 μM) cell assay nor in a genotoxicity assay (no effect in the AMES bacterial reverse mutation assay up to 600 μM). It had minimal liability in the hERG (human ether-a-go-go gene) channel with an IC₅₀ of 47 μM. With the optimal pharmacological, ADME, and pharmacokinetic profiles, compound 7m was scaled up for toxicology and safety studies in rat.

Table 3. In Vivo Pharmacokinetics of 7m^a

	dose (mg/kg) ^g	Cl (mL/min/kg)	V _{ss} (L/kg)	t _{1/2} (h)	oral F (%)
rat ^{b,c}	IV, 1; PO, 5	115 ± 29	8.1 ± 1.2	1.1 ± 0.1	15 ± 7
dog ^{b,d}	IV, 1; PO, 4	21.9	7.3	5.5	67
monkey ^{e,f}	IV, 1; PO, 5	7.6	2.2	6.4	32

^aCl, V_{ss} and t_{1/2} calculations were derived from IV data for all species, and standard error is given for n ≥ 3 measurements. ^b5% DMSO/95% 50 mM lactic acid in saline IV formulation and 0.5% methylcellulose PO formulation. ^cWistar–Han rats (n = 3 animals per dose group). ^dBeagle dogs (n = 2 animals per dose group). ^eCynomolgus monkey (n = 2 animals per dose group). ^f12% sulfobutyl ether beta-cyclodextrin in 50 mM citrate buffer IV formulation and 0.5% methylcellulose PO formulation. ^gIV, intravenous; PO, oral administration.

During the scale up process (beyond 30 g), which involved an acid–base workup, the team discovered that a substantial amount of starting material **6** was regenerated during the workup. Compound **7m** was found to be unstable in both plasma (human and rat) and under aqueous conditions at physiological pH (7.4) and above. The analogous compounds **7j**, **7n**, and **7r** (Table 1) were tested in a pH stability assay at three different pHs (1.5, 6.5, and 7.4). Pyridyl amide **7n** and *N*-methylcarboxamide **7r** were unstable at basic pH. Conversely, compound **7j** was stable under both acidic and basic media. Presumably deprotonation and intramolecular cleavage of the piperidine amide bond by the *ortho*-amide of **7m**, **7n**, and **7r** generates the starting material piperidine **6** (observed by LCMS and ¹H NMR).

It was reasoned that, if the position of the primary *ortho*-amide moiety was moved further away from the piperidine amide, the compounds should be chemically more stable, disrupting the 1,6 relationship between the carbonyl of the piperidine amide and the Lewis basic groups of the pendant carbamide resulting in subsequent piperidine amide cleavage. A set of *ortho*-amides with longer linkers (compounds **7p** and **7o**) were prepared as described above in Scheme 2. These compounds were stable at pH 7.4 but did not meet our potency criteria in the ghrelin binding assay (>100 nM). Interestingly, the nitrile precursor to **7o**, compound **7k**, was one of the most potent compounds in this series with a *K_i* of 4 nM in the binding assay and 5 nM in the functional assay. However, the clearance and MDR ratio did not meet our criteria for advancement. In addition, the reverse *ortho*-amide, **7s**, was prepared from readily available 4-methoxy-2-aminophenylacetic acid and **6** for stability testing purposes. Compound **7s** had reasonable ghrelin binding potency (26 nM) and was stable at pH 7.4, but unfortunately suffered from formation of a reactive metabolite, presumably due to acetamide hydrolysis revealing an aniline that gets bioactivated in liver microsomes.³⁶ It is also worth noting that all compounds **7a–7g** were stable at pH 7.4.

While trying to gain an understanding of the severity of the pH-dependent chemical instability of **7m**, compound **1** progressed to clinical studies and development of compound **7m** was discontinued. In summary, we have described parallel efforts to find a peripherally acting ghrelin inverse agonist via physical chemical property optimization of a centrally acting lead compound **2**. Clear improvements with regards to polarity, potency, clearance, and safety were achieved. Synthesis of polar and heterocyclic piperidine acetamide derivatives delivered several highly potent ghrelin inverse agonists with optimal ADME properties. Compound **7m** was profiled further in various in vitro and in vivo experiments. Chemical instability of the piperidine acetamides bearing an *ortho*-amide functional group was observed. Further optimization of back-up compound **7m** was put on hold pending feedback on the clinical studies with PF-5190457 (**1**).

■ ASSOCIATED CONTENT

■ Supporting Information

Experimental procedures and spectral data. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Notes

The authors declare no competing financial interest.

■ ABBREVIATIONS

HLM, human liver microsome; RLM, rat liver microsome; DLM, dog liver microsome; SFLogD, shake-flask log D; GHS-R1a, growth hormone secretagogue receptor 1a; TPISA, topological polar surface area; MDR, Madin–Darby canine kidney (MDCK) cell line expressing the human MDR-1 gene; EDCI, 1-ethyl-3-(3-(dimethylamino)propyl)carbodiimide; CDI, 1,1'-carbonyldiimidazole; CDMT, 2-chloro-4,6-dimethoxy-1,3,5-triazine; TD, thermodynamic; THLE, transformed human liver epithelial cell assay; hERG, human ether-a-go-go-related gene

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