## Multiple Stable Traveling Wave Profiles of a System of Conservation Laws Arising from Chemotaxis

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**Abstract.** In this paper, we establish the existence and nonlinear stability of a hyperbolic system of conservation laws derived from a repulsive singular chemotaxis model. By the phase plane analysis alongside Poincaré-Bendixson theorem, we first prove that this hyperbolic system admits three different types of traveling wave profiles, which are explicitly illustrated with numerical simulations. Then using a unified weighted energy estimates and technique of taking anti-derivatives, we prove that all types of traveling wave profiles, including non-monotone pulsating wave profiles, are nonlinearly and asymptotically stable if the initial data are small perturbations with zero mass from the spatially shifted traveling wave profiles.

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**Key words**: Chemotaxis, conservation laws, traveling waves, nonlinear stability, weighted energy estimates.

## 1 Introduction

Chemotaxis, the movement of an organism or entity in response to a chemical stimulus, is a widespread phenomenon in nature. One of the pioneering chemotaxis models was proposed by Keller and Segel [17] as follows to describe the wave propagation of bacterial chemotaxis:

$$\begin{cases} u_t = du_{xx} - \chi [u(\ln w)_x]_x, \\ w_t = \varepsilon w_{xx} - \mu u w^m, \end{cases}$$
 (1.1)

where u and w denote the cell density and chemical concentration, respectively. d > 0 and  $\varepsilon \ge 0$  are cell and chemical diffusion coefficients, respectively, and  $m \ge 0$  denotes

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the consumption rate. Generally chemotaxis is said to be attractive if the chemotactic coefficient  $\chi > 0$  and repulsive if  $\chi < 0$  with  $|\chi|$  measuring the strength of chemotaxis. The chemotaxis model (1.1) was used in [17] to describe the process that bacteria u move up (i.e.  $\chi > 0$ ) the concentration gradient of a nutrient (denoted by w) which is absorbed by the bacteria (i.e.  $\mu > 0$ ). A prominent structural feature of (1.1) is that the chemotactic sensitivity function  $\ln w$  is singular at w=0. This imposes tremendous challenges for both analysis and numerical computations. Though the existence/nonexistence of traveling wave solutions for any  $m \geq 0$  have been well understood (cf. [29, 32]), the stability of traveling wave solutions was obtained only for the case m=1 and still widely remains open for the case  $m\neq 1$ . In the case m=1, if  $\chi \mu > 0$ , the following Cole-Hopf transformation (cf. [23, 34]):

$$v = \frac{\sqrt{\chi \mu}}{\mu} \frac{w_x}{w} \tag{1.2}$$

can remove the singularity and convert the system (1.1) into a non-singular system of conservation laws as follows:

$$\begin{cases}
 u_t + (uv)_x = du_{xx}, \\
 v_t + (u + \sigma v^2)_x = \varepsilon v_{xx}
\end{cases}$$
(1.3)

with

$$\sigma = -\frac{\varepsilon}{\chi}$$

where we have used the rescalings  $\tilde{t} = \chi \mu t$  and  $\tilde{x} = \sqrt{\chi \mu} x$  to rescale the model but suppressed the tildes for simplicity. The transformed system (1.3) has no singularity and is more tractable analytically. As  $\chi > 0$  and hence  $\sigma < 0$ , a large amount of interesting results have been developed to the transformed system (1.3) for both  $\varepsilon > 0$  and  $\varepsilon = 0$ , such as traveling wave solutions [1–3, 19–21, 23, 24], global well-posedness of large/small solutions in the whole space [6, 14, 30, 38] or in bounded domains (or intervals) with suitable boundary conditions [22, 31, 40] and boundary layer problem [15]. When  $m \neq 1$ , stability results are restricted only to the spectral stability [25] and absolute instability [5] for the case  $m = \varepsilon = 0$  or instability [28] for  $\varepsilon > 0$ , m = 0. We also refer to a result in [8] for the existence of traveling wave solutions on a generalize Keller-Segel model. Results on the Keller-Segel model (1.1) with fractional diffusion are referred to [12, 13].

The afore-mentioned results are developed for the attractive case  $\chi > 0$ . For the repulsive case  $\chi < 0$ , the Keller-Segel model (1.1) with m = 1 and  $\mu < 0$  was re-derived in [34] based on a random walk framework to describe the biased movement of cells that deposit signals modifying the local environment for subsequent movements, such as myxobacteria or ants that deposit non-diffusive or slowly-moving chemical substances on the way for succeeding passage. In this case, one can still use the Cole-Hopf transformation (1.2) to get (1.3) with  $\sigma = -\varepsilon/\chi > 0$  for which there are also many mathematical results available for the global dynamics in bounded or unbounded domain (cf. [16, 37, 41, 42]). If  $\sigma \ge 0$  is allowed to be an arbitrary constant, the system (1.3) may have more applications. For